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<p>(54) Title: MATERIALS AND METHODS FOR THE MODIFICATION OF PLANT CELL WALL POLYSACCHARIDES</p> <p>(57) Abstract</p> <p>Novel isolated polynucleotides and polypeptides associated with the synthesis of plant cell wall polysaccharides are provided, together with genetic constructs comprising such sequences. Methods for using such constructs for the modulation of polysaccharide content in plants are also disclosed, together with transgenic plants comprising such constructs.</p>		

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## MATERIALS AND METHODS FOR THE MODIFICATION OF PLANT CELL WALL POLYSACCHARIDES

### 5    **Technical Field of the Invention**

This invention relates to the field of modification of cell wall polysaccharide content and composition in plants. More particularly, this invention relates to enzymes involved in the synthesis of plant cell wall polysaccharides and nucleotide sequences encoding such enzymes.

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### **Background of the Invention**

Plant cells are characterised by having a rigid cell wall. These cell walls are comprised primarily of polymers of simple sugar monomers linked in a variety of linear or branched polymers known as polysaccharides. The most abundant simple sugar monomer is glucose, and the most abundant polymer is cellulose. Cellulose is a linear, unbranched polymer, comprised of  $\beta$ -1,4 linked glucose monomers. Other polysaccharides found in plant cell walls include hemicellulose, which is a group of polysaccharides comprised of  $\beta$ -1,4 linked glucose monomers having side chains which may include sugars other than glucose. These side chains frequently include xylose, fucose, arabinose, and galactose. Pectins are another type of polysaccharide found in plant cell walls. Pectins are acidic polysaccharides, which are generally comprised primarily of galacturonic acid and rhamnose sugar monomers. Amylose is an additional common plant polysaccharide which is not usually found as a major component of cell walls. It acts primarily as a storage material for glucose, rather than as a structural polymer. However, because amylose is comprised primarily of  $\alpha$ -1,4-linked glucose monomers, it is considered to be a related polymer from a biochemical and physiological perspective.

Plant polysaccharides have many uses. Certain plastics, such as cellulose acetate, and synthetic textiles, such as rayon, are made from cellulose. In addition, some biodegradable plastics and digestible medicine capsules, as well as medical fillers and fiber additives for food, can be made from plant polysaccharides.

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In foodstuffs, polysaccharides have a profound impact on food quality. Cell walls contribute to crispness in carrots, while degradation of cell walls is required for softening of fruits, such as peaches and tomatoes. In maize, increased amylose is desirable for cattle feed, but not for human consumption, and increased cell wall strength reduces digestibility. In fiber crops, such as timber, cellulose is the primary polymer of interest. Wood density, a fundamental measure of structural timber quality, is essentially a measure of cellulose content. In the paper pulping industry, efficiency is measured in terms of yield of cellulose. Clearly, the ability to increase cellulose content in timber is an important economic goal.

The sugars which make up plant cell wall polysaccharides are produced in the photosynthetic organs of plants. The sugars so produced are commonly converted into sucrose, a disaccharide consisting of glucose and fructose. Sucrose is transported throughout the plant, to wherever sugar monomers are called for. Thus, the photosynthetic organs are often referred to as a source, while tissues requiring large amounts of sugar monomers are referred to as a sink. Actively growing regions of the plant are generally sink tissues, as new cell wall synthesis requires large amounts of sugar monomers.

When the transported sucrose arrives at the sink destination, it must be converted into whichever kind of sugar monomer is required. The sugar monomers which make up plant cell walls are primarily 5- or 6-carbon sugars. Different sugars are generally distinguished by stereospecific orientation of hydroxyl groups. Plants contain a variety of enzymes, such as isomerases or epimerases, which can rapidly change the orientation of these hydroxyls. In addition, there are a number of enzymes which can add or remove a single carbon from a sugar monomer. The result is a single pool of sugar monomers which the plant can freely inter-convert into whichever kind is needed for cell wall synthesis.

Plant polysaccharides are thus biochemically and physiologically inter-related. All polymers compete for the same pool of sugar monomers, and all sugar monomers can be freely interconverted to other types. Degradation of any one polymer will provide building material for any other. Attempts to engineer changes in one polymer may therefore have pleiotropic effects on other polymers.

The rate of cell wall synthesis is dependent on both the availability of sugar monomers to serve as building blocks for the polymers of the wall, and the enzymes which polymerise those building blocks into polymers. Enzymes which are directly responsible for the



synthesis of the major cell wall polymers, such as cellulose, hemicellulose and pectin, may have a profound impact on the rate of cell wall synthesis. Source-sink relations may play an important role in limiting cell wall synthesis, if the availability of substrates becomes limiting. Polymer degrading enzymes may liberate sugar monomers from unnecessary polymers for use  
5 in building new, desired polymers. Enzymes which can isomerise sugars from one form into another can convert the sugars into whichever kind is needed. Each of the different types of cell wall polysaccharides effectively competes for the same pool of sugar monomers, and each represents a potential source of monomers for any of the other polymers.

The final committed steps in cellulose biosynthesis involve a relatively small number  
10 of enzymes. Cellulose synthase (CEL) is believed to function as part of a large, membrane-bound complex which also includes sucrose synthase (SUS: Amor et al., *Proc. Natl. Acad. Sci USA* 92:9353-9357, 1995) and annexin (ANX: Clark and Roux, *Plant Phys.* 109:1133-1139, 1995). This enzyme complex polymerises activated glucose into the cellulose polymer. The glucose is activated by UDP-glucose pyrophosphorylase (UGP), also known as UTP-  
15 glucose-1-phosphate uridylyltransferase. These enzymes are believed to be sufficient for the biosynthesis of cellulose from glucose. Other than these steps, the availability of glucose appears to be the most significant rate-limiting step in cellulose biosynthesis.

Glucose is primarily stored in most plants as amylose. Plants routinely store amylose and degrade it to free up the glucose monomers, as needed. By inhibiting the efficiency of  
20 glucose storage, or by increasing the liberation of glucose from amylose, the availability of glucose monomers for cellulose biosynthesis can be increased. The rate-limiting enzyme in the storage of glucose as amylose is ADP-glucose pyrophosphorylase (AGP), also known as ATP-glucose-1-phosphate adenylyltransferase (Iglesias et al., *J. Biol. Chem.* 268:1081-1086, 1993). Conversely, the enzyme most responsible for liberating glucose from amylose is  
25 amylase (AMA: Kawagoe and Delmer, *Genetic Engineering* 19:63-87, 1997).

These enzymes clearly will be important in the engineering of economically useful changes in cellulose biosynthesis. In addition, there are many other enzymes which may be useful in influencing plant cell wall polysaccharide biosynthesis. Other enzymes likely to be involved in cellulose biosynthesis include 1,4- $\beta$ -cellobiohydrolase,  $\beta$ -glucosidase, calnexin,  
30 cellobiose epimerase, cellobiose phosphorylase, cellulase A, dextranucrase, invertase, phosphodiesterase, phosphoglucomutase, sucrose phosphate synthase, sucrose phosphorylase,

UDP-glucose 4-epimerase and UDP-glucose dehydrogenase. Enzymes believed to be involved in hemicellulose biosynthesis include  $\beta$ -glucanase, arabinan synthase, GDP-fucose pyrophosphorylase, GDP-mannose pyrophosphorylase, 1,3 and 1,4- $\beta$ -glucanases, 1,3 and 1,4- $\beta$ -glucosidases, mannose-6-phosphate isomerase,  $\alpha$ -DP-hexose pyrophosphorylase, xyloglucan endotransglycosylase and xyloglucan synthase. Enzymes likely to be involved in pectin biosynthesis include  $\alpha$ -galactosidase,  $\beta$ -glucuronidase, exopolygalacturonase, glucuronosyl-transferase, pectin methyl-esterase, polygalacturonase and UDP-hexose-1-phosphate uridylyltransferase. Enzymes believed to be involved in amylose biosynthesis include  $\alpha$ -glucosidase, amylopectin 6-glucanohydrolase, amylopectin-branching glycosyltransferase,  $\beta$ -amylase, branching enzyme, inulosucrase, isoamylase, isomaltase, levansucrase, starch phosphorylase and starch synthase. Enzymes likely to be involved in the interconversion of 5-carbon sugars include 2-dehydro-3-deoxy-gluconokinase, aldehyde reductase, arabinose isomerase, D-arabinitol dehydrogenase, D-xylulose reductase, endo-1,4- $\beta$ -xylanase, exo-1,4- $\beta$ -xylanase, L-arabinose isomerase, L-ribulokinase, L-xylulokinase, phospho-ribulokinase, ribose 5-phosphate isomerase, ribulose-phosphate-3-epimerase, ribulose-phosphate-4-epimerase, transaldolase, transketolase, xylose isomerase and xylulokinase. Enzymes likely to be involved in interconversion of 6-carbon sugars include 6-phospho-fructo-1-kinase, 6-phospho-fructo-2-kinase, trehalose phosphate synthase, aldolase, aldose 1-epimerase, D-fructokinase, D-galactokinase, fructose 1,6-diphosphatase, gluconolactonase, glucose 1-phosphatase, glucose 6-phosphatase, glucose 6-phosphate dehydrogenase, glucose-phosphate isomerase, hexokinase, phosphoglucomutase, trehalase, trehalose phosphatase and UDP-galactose dehydrogenase.

While DNA sequences encoding some of the enzymes involved in the biosynthetic pathways of plant cell wall polysaccharides have been isolated for certain species of plants, genes encoding many of the enzymes in a wide range of plant species have not yet been identified. Thus, there remains a need in the art for materials useful in the modification of cell wall polysaccharide content and composition in plants.

### Summary of the Invention

Briefly, the present invention provides polynucleotides isolated from eucalyptus and pine which encode enzymes involved in the synthesis of cell wall polysaccharides. Genetic constructs including such sequences and methods for the use of such constructs are also provided, together with transgenic plants having altered cell wall polysaccharide content and composition.

In one embodiment, the isolated polynucleotides comprise a nucleotide sequence selected from the group consisting of: (a) sequences recited in SEQ ID NOS: 1-29, 57-80, 105, 107, 109-113, 119-129, 139-143 and 149-908; (b) complements of the sequences recited in SEQ ID NOS: 1-29, 57-80, 105, 107, 109-113, 119-129, 139-143 and 149-908; (c) reverse complements of the sequences recited in SEQ ID NOS: 1-29, 57-80, 105, 107, 109-113, 119-129, 139-143 and 149-908; (d) reverse sequences of the sequences recited in SEQ ID NOS: 1-29, 57-80, 105, 107, 109-113, 119-129, 139-143 and 149-908; and (e) sequences having either 40%, 60%, 75% or 90% identical nucleotides, as defined herein, to a sequence of (a) - (d).

In a further aspect, isolated polypeptides encoded by a polynucleotide of the present invention are provided. In one embodiment, such polypeptides comprise an amino acid sequence selected from the group consisting of SEQ ID NOS: 30-56, 81-104, 106, 108, 114-118, 129-138 and 144-148, and variants thereof.

In another aspect, the invention provides genetic constructs comprising a polynucleotide of the present invention, either alone, in combination with one or more of the inventive polynucleotide sequences, or in combination with one or more known polynucleotides, together with transgenic cells comprising such constructs.

In a related aspect, the present invention provides genetic constructs comprising, in the 5'-3' direction, a gene promoter sequence; an open reading frame coding for at least a functional portion of an enzyme encoded by a polynucleotide of the present invention or a variant thereof; and a gene termination sequence. The open reading frame may be orientated in either a sense or antisense direction. Genetic constructs comprising a non-coding region of a gene coding for an enzyme encoded by the above polynucleotides or a nucleotide sequence complementary to a non-coding region, together with a gene promoter sequence and a gene termination sequence, are also provided. Preferably, the gene promoter and termination

sequences are functional in a host plant. Most preferably, the gene promoter and termination sequences are those of the original enzyme genes but others generally used in the art, such as the Cauliflower Mosaic Virus (CMV) promoter, with or without enhancers such as the Kozak sequence or Omega enhancer, and *Agrobacterium tumefaciens* nopal synthase terminator  
5 may be usefully employed in the present invention. Tissue-specific promoters may be employed in order to target expression to one or more desired tissues. In a preferred embodiment, the gene promoter sequence provides for transcription in xylem. The genetic construct may further include a marker for the identification of transformed cells.

In a further aspect, transgenic plant cells comprising the genetic constructs of the  
10 present invention are provided, together with plants comprising such transgenic cells, and fruits, seeds and other products, derivatives, or progeny of such forestry plants. Propagules of the transgenic plants transformed with the inventive polynucleotides are also included in the present invention. As used herein, the word "propagule" means any part of a plant that may be used in reproduction or propagation, sexual or asexual, including cuttings.

15 Plant varieties, particularly registrable plant varieties according to Plant Breeders' Rights, may be excluded from the present invention. A plant need not be considered a "plant variety" simply because it contains stably within its genome a transgene, introduced into a cell of the plant or an ancestor thereof.

In yet another aspect, methods for modulating the polysaccharide content and  
20 composition of an organism, such as a plant, are provided, such methods including stably incorporating into the genome of the plant a genetic construct of the present invention. In a preferred embodiment, the target plant is a woody plant, preferably selected from the group consisting of eucalyptus, pine, acacia, poplar, sweetgum, teak and mahogany species, more preferably from the group consisting of pine and eucalyptus species, and most preferably from  
25 the group consisting of *Eucalyptus grandis* and *Pinus radiata*. In a related aspect, a method for producing a plant having modified cellulose content is provided, the method comprising transforming a plant cell with a genetic construct of the present invention to provide a transgenic cell and cultivating the transgenic cell under conditions conducive to regeneration and mature plant growth.

30 In yet a further aspect, the present invention provides methods for modifying the activity of a polypeptide in a plant, comprising stably incorporating into the genome of the

plant a genetic construct of the present invention. In a preferred embodiment, the target plant is a woody plant, preferably selected from the group consisting of eucalyptus, pine, acacia, poplar, sweetgum, teak and mahogany species, more preferably from the group consisting of pine and eucalyptus species, and most preferably from the group consisting of *Eucalyptus grandis* and *Pinus radiata*.

The above-mentioned and additional features of the present invention and the manner of obtaining them will become apparent, and the invention will be best understood by reference to the following more detailed description. All references disclosed herein are hereby incorporated by reference in their entirety as if each was incorporated individually.

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#### Brief Description of the Figures

Fig. 1 illustrates the level of native CEL enzyme activity in positive control mung bean (*V. radiata*) plants.

Fig. 2 illustrates the level of CEL enzyme activity in mammalian 293T cells transfected with *E. grandis* CEL as compared to that in non-transfected 293T cells.

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#### Detailed Description

As outlined above, cellulose is formed by polymerization of glucose into a linear, unbranched, polymer comprised of  $\beta$ -1,4 linked glucose monomers (Kawagoe and Delmer, *Genetic Engineering*, 19:63-87, 1997). Cellulose is the most important plant cell wall polysaccharide from both a structural, as well as industrial, perspective. Other polysaccharides are essential for healthy cell walls, as well as for many alternative industrial uses.

Glucose monomers are most commonly stored in the plant in the form of amylose by the action of several enzymes, with the rate limiting step for storage being catalysed by AGP (Iglesias et al., *J. Biol. Chem.* 268:1081-1086). Glucose monomers are freed from this storage polymer by the action of the enzyme AMA. The free monomers are activated by the action of the enzyme UGP, and polymerised into cellulose macro-crystalline structures by the action of the cellulose synthase enzyme complex. Pure CEL enzyme has been shown to form  $\beta$ -1,4 glucose linkages *in vitro*, but has not been shown to be sufficient for polymerization of the

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large polymers which are fundamental to the structure of plant cell walls. The holoenzyme complex appears to be necessary for this latter function. The holoenzyme is believed to be comprised of the CEL enzyme in combination with the SUS enzyme and ANX, the whole complex being integrated into the plasma membrane and forming a "rosette" structure as seen  
5 in electron micrographs of plant cell membranes (Arioli et al., *Science* 279:717-720, 1998).

Because cellulose synthesis can represent such a large sink for sugar monomers in the cell, changes in the rate of cellulose synthesis can have a profound influence on the synthesis of other plant polysaccharides. Conversely, changes in the rates of synthesis of other plant polysaccharides can have a profound influence on the pool of sugars available for synthesis of  
10 cellulose. Hence, changes in the synthesis of any single polymer may affect both the content and composition of plant cell wall polysaccharides, and polysaccharides in general.

Quantitative and qualitative modifications in plant polysaccharide content are known to be induced by external factors such as light stimulation, low calcium levels, and mechanical stress. Synthesis of cell wall polysaccharides can also be induced by infection  
15 with pathogens.

Using the methods and materials of the present invention, the polysaccharide content of a plant may be increased or reduced, by incorporating additional copies of genes encoding enzymes involved in the synthesis of cell wall polysaccharides into the genome of the target plant. Similarly, an increase or decrease in polysaccharide content may be obtained by  
20 transforming the target plant with antisense copies of such genes. In addition, the number of copies of genes encoding for different enzymes in the biosynthetic pathway of cell wall polysaccharides can be manipulated to modify the relative amount of each monosaccharide synthesized, thereby leading to the formation of cell walls having altered composition. The alteration of polysaccharide composition would be advantageous, for example, in tree  
25 processing for paper.

The polynucleotides of the present invention were isolated from forestry plant sources, namely from *Eucalyptus grandis* and *Pinus radiata*, but they may alternatively be synthesized using conventional synthesis techniques. Specifically, isolated polynucleotides of the present invention include polynucleotides comprising a sequence selected from the group consisting  
30 of sequences identified as SEQ ID NOS: 1-29, 57-80, 105, 107, 109-113, 119-129, 139-143 and 149-908; complements of the sequences identified as SEQ ID NOS: 1-29, 57-80, 105,

107, 109-113, 119-129, 139-143 and 149-908; reverse complements of the sequences identified as SEQ ID NOS: 1-29, 57-80, 105, 107, 109-113, 119-129, 139-143 and 149-908; at least a specified number of contiguous residues (x-mers) of any of the above-mentioned polynucleotides; extended sequences corresponding to any of the above polynucleotides; antisense sequences corresponding to any of the above polynucleotides; and variants of any of the above polynucleotides, as that term is described in this specification.

In another embodiment, the present invention provides isolated polypeptides encoded by the DNA sequences of SEQ ID NOS: 1-29, 57-80, 105, 107, 109-113, 119-129, 139-143 and 149-908;. The predicted amino acid sequences encoded by SEQ ID NOS: 1-22, 24-28, 57-80, 105, 107, 109-113 and 119-143, based on the best available information at the time of filing this application, are provided in SEQ ID NOS: 30-56, 81-104, 106, 108, 114-118, 129-138 and 144-148, respectively. The present invention also encompasses polynucleotides that differ from the disclosed sequences but which, due to the degeneracy of the genetic code, encode a polypeptide which is the same as that encoded by a polypeptide of the present invention. Such polynucleotides are said to be "degeneratively equivalent" to a polynucleotide sequence disclosed herein.

The polynucleotides and polypeptides of the present invention were putatively identified by DNA and polypeptide similarity searches. In the attached Sequence Listing SEQ ID NOS: 1-29, 57-80, 105, 107, 109-113, 119-129, 139-143 and 149-908 are polynucleotide sequences, and SEQ ID NOS: 30-56, 81-104, 106, 108, 114-118, 129-138 and 144-148 are polypeptide sequences. The polynucleotides and polypeptides of the present invention, have demonstrated similarity to enzymes that are known to be involved in the synthesis of cell wall polysaccharides. The putative identity of each of the inventive polynucleotides is shown below in Table 1.

TABLE 1

DNA SEQ ID NO:	PROTEIN SEQ ID NO:	IDENTITY
1	30	AGP
2	31	AGP
3	32	AGP
4	33	AMA

DNA SEQ ID NO:	PROTEIN SEQ ID NO:	IDENTITY
5	34	AMA
6	35	AMA
7	36	CEL
8	37	CEL
9	38	CEL
10	39	CEL
11	40	CEL
12	41	CEL
13	42	CEL
14	43	CEL
15	44	SUS
16	45	SUS
17	46	SUS
18	47	SUS
19	48	SUS
20	49	UGP
21	50	UGP
22	51	UGP
23	-	UGP
24	52	ANX
25	53	ANX
26	54	ANX
27	55	ANX
28	56	ANX
29	-	ANX
57	81	AMA
58	82	AMA
59	83	AGP
60	84	AGP
61	85	AGP
62	86	AGP
63	87	AGP
64	88	AGP
65	89	AGP
66	90	CEL
67	91	CEL
68	92	CEL
69	93	CEL
70	94	CEL
71	95	SUS
72	96	SUS
73	97	SUS
74	98	SUS
75	99	SUS
76	100	SUS
77	101	SUS
78	102	SUS
79	103	UGP
80	104	UGP



DNA SEQ ID NO:	PROTEIN SEQ ID NO:	IDENTITY
105	106	SUS
107	108	CEL
109	114	ANX
110	115	ANX
111	116	ANX
112	117	ANX
113	118	ANX
119	129	CEL
120	130	CEL
121	131	CEL
122	132	CEL
123	133	CEL
124	134	CEL
125	135	CEL
126	136	CEL
127	137	CEL
128	138	CEL
135	144	SUS
140	145	$\alpha$ -amylase
141	146	CEL
142	147	AGP (3' end of SEQ ID NO: 62)
143	148	SUS (3' of SEQ ID NO: 74)
149-185	-	1,3- $\beta$ -D-Glucanase
186	-	1,4- $\beta$ -Cellobiohydrolase
187-196	-	$\alpha,\alpha$ -trehalose phosphate synthase
197-204	-	$\alpha$ -glucosidase
205-250	-	aldolase
251	-	Amylopectin 6-glucanohydrolase
252-262	-	$\beta$ -amylase
263	-	$\beta$ -glucosidase
264-272	-	Branching enzyme
273-318	-	D-fructokinase
319-354	-	D-xylulose reductase
355-365	-	Endo-1,3-1,4- $\beta$ -glucanase
366-371	-	Glucan exo-1,3- $\beta$ -glucosidase
372-377	-	Glucose 6-phosphate dehydrogenase
378-381	-	Glucose phosphate isomerase
382-389	-	Isoamylase
390-393	-	L-ribulokinase
394-398	-	Mannitol-1-phosphate 5-dehydrogenase
399-478	-	Pectin methyl-esterase
479-506	-	Phosphoglucomutase
507-508	-	Phospho-ribulokinase
509-521	-	Ribulose-phosphate-3-epimerase
522-530	-	Starch phosphorylase
531-551	-	Sucrose phosphate synthase
552-555	-	SUS
556-586	-	Transketolase
587-591	-	Trehalase

DNA SEQ ID NO:	PROTEIN SEQ ID NO:	IDENTITY
592-620	-	UDP-glucose 4-epimerase
621-902	-	Xyloglucan endotransglycosylase
903-908	-	Xylose isomerase

The term "polynucleotide(s)," as used herein, means a single or double-stranded polymer of deoxyribonucleotide or ribonucleotide bases and includes DNA and corresponding RNA molecules, including HnRNA and mRNA molecules, both sense and anti-sense strands, and comprehends cDNA, genomic DNA and recombinant DNA, as well as wholly or partially synthesized polynucleotides. An HnRNA molecule contains introns and corresponds to a DNA molecule in a generally one-to-one manner. An mRNA molecule corresponds to an HnRNA and DNA molecule from which the introns have been excised. A polynucleotide may consist of an entire gene, or any portion thereof. Operable anti-sense polynucleotides may comprise a fragment of the corresponding polynucleotide, and the definition of "polynucleotide" therefore includes all such operable anti-sense fragments.

The term "polypeptide", as used herein, encompasses amino acid chains of any length including full length proteins, wherein amino acid residues are linked by covalent peptide bonds. Polypeptides of the present invention may be naturally purified products, or may be produced partially or wholly using recombinant techniques.

The definition of the terms "complement", "reverse complement" and "reverse sequence", as used herein, is best illustrated by the following example. For the sequence 5' AGGACC 3', the complement, reverse complement and reverse sequence are as follows:

complement	3' TCCTGG 5'
reverse complement	3' GGTCCT 5'
reverse sequence	5' CCAGGA 3'.

As used herein, the term "variant" covers any sequence which has at least about 40%, more preferably at least about 60%, more preferably yet at least about 75% and most preferably at least about 90% identical residues (either nucleotides or amino acids) to a sequence of the present invention. The percentage of identical residues is determined by aligning the two sequences to be compared, determining the number of identical residues in

the aligned portion, dividing that number by the total length of the inventive, or queried, sequence and multiplying the result by 100.

Polynucleotide or polypeptide sequences may be aligned, and percentage of identical nucleotides in a specified region may be determined against another polynucleotide, using  
5 computer algorithms that are publicly available. Two exemplary algorithms for aligning and identifying the similarity of polynucleotide sequences are the BLASTN and FASTA algorithms. The similarity of polypeptide sequences may be examined using the BLASTP algorithm. Both the BLASTN and BLASTP software are available on the NCBI anonymous FTP server (<ftp://ncbi.nlm.nih.gov>) under /blast/executables/. The BLASTN algorithm  
10 Version 2.0.6 [Sept-16-1998], set to the default parameters described in the documentation and distributed with the algorithm, is preferred for use in the determination of variants according to the present invention. The use of the BLAST family of algorithms, including BLASTN and BLASTP, is described at NCBI's Internet website at the URL <http://www.ncbi.nlm.nih.gov/BLAST/newblast.html> and in the publication of Altschul,  
15 Stephen F, et al., "Gapped BLAST and PSI-BLAST: a new generation of protein database search programs," *Nucleic Acids Res.* 25:3389-3402, 1997. The computer algorithm FASTA is available on the Internet at the ftp site <ftp://ftp.virginia.edu/pub/fasta/>. Version 2.04, [February 1996], set to the default parameters described in the documentation and distributed with the algorithm, is preferred for use in the determination of variants according to the  
20 present invention. The use of the FASTA algorithm is described in Pearson WR and Lipman DJ, "Improved Tools for Biological Sequence Analysis," *Proc. Natl. Acad. Sci. USA* 85:2444-2448, 1988; and Pearson WR, "Rapid and Sensitive Sequence Comparison with FASTP and FASTA," *Methods in Enzymol.* 183:63-98, 1990.

The following running parameters are preferred for determination of alignments and  
25 identities using BLASTN that contribute to the E values and percentage identity of polynucleotides of the present invention: Unix running command: blastall -p blastn -d embldb -e 10 -G 0 -E 0 -r 1 -v 30 -b 30 -i queryseq -o results; and the parameters are: -p Program Name [String]; -d Database [String]; -e Expectation value (E) [Real]; -G Cost to open a gap (zero invokes default behavior) [Integer]; -E Cost to extend a gap (zero invokes default  
30 behavior) [Integer]; -r Reward for a nucleotide match (blastn only) [Integer]; -v Number of

one-line descriptions (V) [Integer]; -b Number of alignments to show (B) [Integer]; -i Query File [File In]; -o BLAST report Output File [File Out] Optional.

The following running parameters are preferred for determination of alignments and identities using BLASTP that contribute to the E values and percentage identity of polypeptide sequences: For BLASTP the following running parameters are preferred: blastall -p blastp -d swissprot -e 10 -G 0 -E 0 -v 30 -b 30 -i queryseq -o results; and parameters are: -p Program Name [String]; -d Database [String]; -e Expectation value (E) [Real]; -G Cost to open a gap (zero invokes default behavior) [Integer]; -E Cost to extend a gap (zero invokes default behavior) [Integer]; -v Number of one-line descriptions (v) [Integer]; -b Number of alignments to show (b) [Integer]; -I Query File [File In]; -o BLAST report Output File [File Out] Optional.

The "hits" to one or more database sequences by a queried sequence produced by BLASTN, BLASTP, FASTA, or a similar algorithm, align and identify similar portions of sequences. The hits are arranged in order of the degree of similarity and the length of sequence overlap. Hits to a database sequence generally represent an overlap over only a fraction of the sequence length of the queried sequence.

The BLASTN and FASTA algorithms also produce "Expect" values for alignments. The Expect value (E) indicates the number of hits one can "expect" to see over a certain number of contiguous sequences by chance when searching a database of a certain size. The Expect value is used as a significance threshold for determining whether the hit to a database, such as the preferred EMBL database, indicates true similarity. For example, an E value of 0.1 assigned to a hit is interpreted as meaning that in a database of the size of the EMBL database, one might expect to see 0.1 matches over the aligned portion of the sequence with a similar score simply by chance. By this criterion, the aligned and matched portions of the sequences then have a probability of 90% of being the same. For sequences having an E value of 0.01 or less over aligned and matched portions, the probability of finding a match by chance in the EMBL database is 1% or less using the BLASTN or FASTA algorithm.

According to one embodiment, "variant" polynucleotides, with reference to each of the polynucleotides of the present invention, preferably comprise sequences having the same number or fewer nucleic acids than each of the polynucleotides of the present invention and producing an E value of 0.01 or less when compared to the polynucleotide of the present

invention. That is, a variant polynucleotide is any sequence that has at least a 99% probability of being the same as the polynucleotide of the present invention, measured as having an E value of 0.01 or less using the BLASTN or FASTA algorithms set at the default parameters. According to a preferred embodiment, a variant polynucleotide is a sequence having the same  
5 number or fewer nucleic acids than a polynucleotide of the present invention that has at least a 99% probability of being the same as the polynucleotide of the present invention, measured as having an E value of 0.01 or less using the BLASTN or FASTA algorithms set at the default parameters.

Alternatively, variant polynucleotide hybridize to the polynucleotide of the present  
10 invention under stringent conditions. As used herein, "stringent conditions" refers to prewashing in a solution of 6X SSC, 0.2% SDS; hybridizing at 65°C, 6X SSC, 0.2% SDS overnight; followed by two washes of 30 minutes each in 1X SSC, 0.1% SDS at 65°C and two washes of 30 minutes each in 0.2X SSC, 0.1% SDS at 65°C.

The present invention also encompasses polynucleotides that differ from the disclosed  
15 sequences but that, as a consequence of the discrepancy of the genetic code, encode a polypeptide having similar enzymatic activity as a polypeptide encoded by a polynucleotide of the present invention. Thus, polynucleotides comprising sequences that differ from the polynucleotide sequences recited in SEQ ID NOS: 1-29, 57-80, 105, 107, 109-113, 119-129, 139-143 and 149-908, or complements, reverse sequences, or reverse complements of those  
20 sequences as a result of conservative substitutions are contemplated by and encompassed within the present invention. Additionally, polynucleotides comprising sequences that differ from the polynucleotide sequences recited in SEQ ID NOS: 1-29, 57-80, 105, 107, 109-113, 119-129, 139-143 and 149-908, or complements, reverse complements, or reverse sequences as a result of deletions and/or insertions totaling less than 10% of the total sequence length are  
25 also contemplated by and encompassed within the present invention. Similarly, polypeptides comprising sequences that differ from the polypeptide sequences recited in SEQ ID NOS: 30-56, 81-104, 106, 108, 114-118, 129-138 and 144-148 as a result of amino acid substitutions, insertions, and/or deletions totaling less than 10% of the total sequence length are contemplated by an encompassed within the present invention, provided the variant  
30 polypeptide has activity in a cell wall polysaccharide synthesis pathway.

1 Variants of the polypeptide sequences recited in SEQ ID NOS: 30-56, 81-104, 106,  
108, 114-118, 129-138 and 144-148, wherein the variant has an activity level that is different  
to that of the recited polypeptide are also encompassed by the present invention. In specific  
embodiments, variants of the inventive sucrose synthase (SUS) polypeptides are provided  
5 wherein the N-terminal serine phosphorylation site has been replaced by an acidic amino acid  
(such as Asp or Glu) by, for example, site directed mutagenesis. Nakai et al. have  
demonstrated that SUS polypeptides mutated in this manner possess increased activity  
compared to wild-type SUS (Nakai et al., *Plant Cell Physiol.* 39:1337-1341, 1998).  
Polynucleotides encoding such variants of the inventive SUS polypeptides may therefore be  
10 employed in transgenic plants to increase cellulose production.

The polynucleotides of the present invention may be isolated from various libraries, or  
may be synthesized using techniques that are well known in the art. The polynucleotides may  
be synthesized, for example, using automated oligonucleotide synthesizers (e.g., Beckman  
Oligo 1000M DNA Synthesizer) to obtain polynucleotide segments of up to 50 or more  
15 nucleic acids. A plurality of such polynucleotide segments may then be ligated using standard  
DNA manipulation techniques that are well known in the art of molecular biology. One  
conventional and exemplary polynucleotide synthesis technique involves synthesis of a single  
stranded polynucleotide segment having, for example, 80 nucleic acids, and hybridizing that  
segment to a synthesized complementary 85 nucleic acid segment to produce a 5 nucleotide  
20 overhang. The next segment may then be synthesized in a similar fashion, with a 5 nucleotide  
overhang on the opposite strand. The "sticky" ends ensure proper ligation when the two  
portions are hybridized. In this way, a complete polynucleotide of the present invention may  
be synthesized entirely *in vitro*.

Some of the polynucleotides identified as SEQ ID NOS: 1-29, 57-80, 105, 107,  
25 109-113, 119-129, 139-143 and 149-908 are referred to as "partial" sequences, in that they do  
not represent the full coding portion of a gene encoding a naturally occurring polypeptide.  
The partial polynucleotide sequences disclosed herein may be employed to obtain the  
corresponding full length genes for various species and organisms by, for example, screening  
DNA expression libraries using hybridization probes based on the polynucleotides of the  
30 present invention, or using PCR amplification with primers based upon the polynucleotides of  
the present invention. In this way one can, using methods well known in the art, extend a

polynucleotide of the present invention upstream and downstream of the corresponding mRNA, as well as identify the corresponding genomic DNA, including the promoter and enhancer regions, of the complete gene. The present invention thus comprehends isolated polynucleotides comprising a sequence identified in SEQ ID NOS: 1-29, 57-80, 105, 107, 109-113, 119-129, 139-143 and 149-908, or a variant of one of the specified sequences, that encode a functional polypeptide, including full length genes. Such extended polynucleotides may have a length of from about 50 to about 4,000 nucleic acids or base pairs, and preferably have a length of less than about 4,000 nucleic acids or base pairs, more preferably yet a length of less than about 3,000 nucleic acids or base pairs, more preferably yet a length of less than about 2,000 nucleic acids or base pairs. Under some circumstances, extended polynucleotides of the present invention may have a length of less than about 1,800 nucleic acids or base pairs, preferably less than about 1,600 nucleic acids or base pairs, more preferably less than about 1,400 nucleic acids or base pairs, more preferably yet less than about 1,200 nucleic acids or base pairs, and most preferably less than about 1,000 nucleic acids or base pairs.

Polynucleotides of the present invention also comprehend polynucleotides comprising at least a specified number of contiguous residues ( $x$ -mers) of any of the polynucleotides identified as SEQ ID NOS: 1-29, 57-80, 105, 107, 109-113, 119-129, 139-143 and 149-908, complements, reverse sequences, and reverse complements of such sequences, and their variants. Similarly, polypeptides of the present invention comprehend polypeptides comprising at least a specified number of contiguous residues ( $x$ -mers) of any of the polypeptides identified as SEQ ID NOS: 30-56, 81-104, 106, 108, 114-118, 129-138 and 144-148, and their variants. As used herein, the term " $x$ -mer," with reference to a specific value of " $x$ ," refers to a sequence comprising at least a specified number (" $x$ ") of contiguous residues of any of the polynucleotides identified as SEQ ID NO: 1-29, 57-80, 105, 107, 109-113, 119-129, 139-143 and 149-908, or the polypeptides identified as SEQ ID NOS: 30-56, 81-104, 106, 108, 114-118, 129-138 and 144-148. According to preferred embodiments, the value of  $x$  is preferably at least 20; more preferably, at least 40; more preferably yet, at least 60; and most preferably, at least 80. Thus, polynucleotides and polypeptides of the present invention comprise a 20-mer, a 40-mer, a 60-mer, an 80-mer, a 100-mer, a 120-mer, a 150-mer, a 180-mer, a 220-mer, a 250-mer, or a 300-mer, 400-mer,

500-mer or 600-mer of a polynucleotide or polypeptide identified as SEQ ID NOS: 1-908, and variants thereof.

Polynucleotide probes and primers complementary to and/or corresponding to SEQ ID NOS: 1-29, 57-80, 105, 107, 109-113, 119-129, 139-143 and 149-908, and variants of those  
5 sequences, are also comprehended by the present invention. Such oligonucleotide probes and primers are substantially complementary to the polynucleotide of interest. As used herein, the term "oligonucleotide" refers to a relatively short segment of a polynucleotide sequence, generally comprising between 6 and 60 nucleotides, and comprehends both probes for use in hybridization assays and primers for use in the amplification of DNA by polymerase chain  
10 reaction.

An oligonucleotide probe or primer is described as "corresponding to" a polynucleotide of the present invention, including one of the sequences set out as SEQ ID NOS: 1-29, 57-80, 105, 107, 109-113, 119-129, 139-143 and 149-908, or a variant, if the oligonucleotide probe or primer, or its complement, is contained within one of the sequences  
15 set out as SEQ ID NOS: 1-29, 57-80, 105, 107, 109-113, 119-129, 139-143 and 149-908, or a variant of one of the specified sequences.

Two single stranded sequences are said to be substantially complementary when the nucleotides of one strand, optimally aligned and compared, with the appropriate nucleotide insertions and/or deletions, pair with at least 80%, preferably at least 90% to 95%, and more  
20 preferably at least 98% to 100%, of the nucleotides of the other strand. Alternatively, substantial complementarity exists when a first DNA strand will selectively hybridize to a second DNA strand under stringent hybridization conditions. Stringent hybridization conditions for determining complementarity include salt conditions of less than about 1 M, more usually less than about 500 mM, and preferably less than about 200 mM. Hybridization  
25 temperatures can be as low as 5°C, but are generally greater than about 22°C, more preferably greater than about 30°C, and most preferably greater than about 37°C. Longer DNA fragments may require higher hybridization temperatures for specific hybridization. Since the stringency of hybridization may be affected by other factors such as probe composition, presence of organic solvents and extent of base mismatching, the combination of parameters  
30 is more important than the absolute measure of any one alone. The DNA from plants or



samples or products containing plant material can be either genomic DNA or DNA derived by preparing cDNA from the RNA present in the sample.

In addition to DNA-DNA hybridization, DNA-RNA or RNA-RNA hybridization assays are also possible. In the first case, the mRNA from expressed genes would then be  
5 detected instead of genomic DNA or cDNA derived from mRNA of the sample. In the second case, RNA probes could be used. In addition, artificial analogs of DNA hybridizing specifically to target sequences could also be used.

In specific embodiments, the oligonucleotide probes and/or primers comprise at least about 6 contiguous residues, more preferably at least about 10 contiguous residues, and most  
10 preferably at least about 20 contiguous residues complementary to a polynucleotide sequence of the present invention. Probes and primers of the present invention may be from about 8 to 100 base pairs in length or, preferably from about 10 to 50 base pairs in length or, more preferably from about 15 to 40 base pairs in length. The probes can be easily selected using procedures well known in the art, taking into account DNA-DNA hybridization stringencies,  
15 annealing and melting temperatures, and potential for formation of loops and other factors, which are well known in the art. Tools and software suitable for designing probes, and especially suitable for designing PCR primers, are available on the Internet, for example, at URL <http://www.horizonpress.com/pcr/>. Preferred techniques for designing PCR primers are also disclosed in Dieffenbach CW and Dykster GS, *PCR primer: a laboratory manual*,  
20 CSHL Press: Cold Spring Harbor, NY, 1995.

A plurality of oligonucleotide probes or primers corresponding to a polynucleotide of the present invention may be provided in a kit form. Such kits generally comprise multiple DNA or oligonucleotide probes, each probe being specific for a polynucleotide sequence. Kits of the present invention may comprise one or more probes or primers corresponding to a  
25 polynucleotide of the present invention, including a polynucleotide sequence identified in SEQ ID NOS: 1-29, 57-80, 105, 107, 109-113, 119-129, 139-143 and 149-908.

In one embodiment, the present invention provides genetic that include an open reading frame coding for at least a functional portion of a polypeptide encoded by a polynucleotide of the present invention or a variant thereof. As used herein, the "functional  
30 portion" of a polypeptide is that portion which contains the active site essential for affecting the metabolic step, *i.e.*, the portion of the molecule that is capable of binding one or more

reactants or is capable of improving or regulating the rate of reaction. The functional portion can be determined by targeted mutagenesis and screening of modified protein products with protocols well known in the art. Normally, the functional portion is 10-20 amino acids, but can be shorter or longer. The active site may be made up of separate portions present on one or more polypeptide chains and will generally exhibit high substrate specificity. The term "polypeptide encoded by a polynucleotide" as used herein, includes polypeptides encoded by a nucleotide sequence which includes the partial isolated DNA sequences of the present invention.

The open reading frame may be inserted in the genetic construct in a sense or antisense orientation, such that transformation of a target plant with the genetic construct will produce a change in the amount or structure of the polypeptide compared to the wild-type plant. Transformation with a genetic construct comprising an open reading frame in a sense orientation will generally result in modified expression of the selected gene, while transformation with a genetic construct comprising an open reading frame in an antisense orientation also generally results in modified expression of the selected gene. A population of plants transformed with a genetic construct comprising an open reading frame of the present invention in either a sense or antisense orientation may be screened for increased or reduced expression of the gene in question using techniques well known to those of skill in the art, and plants having the desired phenotypes may thus be isolated.

Alternatively, expression of a gene involved in the biosynthesis of polysaccharides may be inhibited by inserting a portion of an open reading frame of the present invention, in either sense or antisense orientation, in the genetic construct. Such portions need not be full-length but preferably comprise at least 25 and more preferably at least 50 residues of a polynucleotide of the present invention. A much longer portion or even the full length polynucleotide corresponding to the complete open reading frame may be employed. The portion of the open reading frame does not need to be precisely the same as the endogenous sequence, provided that there is sufficient sequence similarity to achieve inhibition of the target gene. Thus a sequence derived from one species may be used to inhibit expression of a gene in a different species.

In a second embodiment, the inventive genetic constructs comprise a polynucleotide including a non-coding region of a gene coding for a polypeptide encoded by a polynucleotide



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<b>(21) International Application Number:</b> PCT/NZ99/00169  <b>(22) International Filing Date:</b> 8 October 1999 (08.10.99)  <b>(30) Priority Data:</b> 09/170,862      13 October 1998 (13.10.98)      US 60/148,426      11 August 1999 (11.08.99)      US  <b>(71) Applicants (for all designated States except US):</b> GENESIS RE- SEARCH AND DEVELOPMENT CORPORATION LIM- ITED [NZ/NZ]; 1 Fox Street, Parnell, Auckland (NZ). FLETCHER CHALLENGE FORESTS LIMITED [NZ/NZ]; 585 Great South Road, Penrose, Auckland (NZ).  <b>(72) Inventor; and</b> <b>(75) Inventor/Applicant (for US only):</b> BLOKSBERG, Leonard, Nathan [US/NZ]; 5A Korau Road, Greenlane, Auckland (NZ).  <b>(74) Agents:</b> BENNETT, Michael, Roy et al.; West-Walker Ben- nett, Mobil on the Park, 157 Lambton Quay, Wellington (NZ).		<b>(81) Designated States:</b> AE, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CR, CU, CZ, DE, DK, DM, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, US, UZ, VN, YU, ZA, ZW, ARIPO patent (GH, GM, KE, LS, MW, SD, SL, SZ, TZ, UG, ZW), Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European patent (AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE), OAPI patent (BF, BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG).  <b>Published</b> <i>Without international search report and to be republished          upon receipt of that report.</i>
<b>(54) Title:</b> MATERIALS AND METHODS FOR THE MODIFICATION OF PLANT CELL WALL POLYSACCHARIDES  <b>(57) Abstract</b>  Novel isolated polynucleotides and polypeptides associated with the synthesis of plant cell wall polysaccharides are provided, together with genetic constructs comprising such sequences. Methods for using such constructs for the modulation of polysaccharide content in plants are also disclosed, together with transgenic plants comprising such constructs.		

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## MATERIALS AND METHODS FOR THE MODIFICATION OF PLANT CELL WALL POLYSACCHARIDES

### 5    **Technical Field of the Invention**

This invention relates to the field of modification of cell wall polysaccharide content and composition in plants. More particularly, this invention relates to enzymes involved in the synthesis of plant cell wall polysaccharides and nucleotide sequences encoding such enzymes.

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### **Background of the Invention**

Plant cells are characterised by having a rigid cell wall. These cell walls are comprised primarily of polymers of simple sugar monomers linked in a variety of linear or branched polymers known as polysaccharides. The most abundant simple sugar monomer is glucose, and the most abundant polymer is cellulose. Cellulose is a linear, unbranched polymer, comprised of  $\beta$ -1,4 linked glucose monomers. Other polysaccharides found in plant cell walls include hemicellulose, which is a group of polysaccharides comprised of  $\beta$ -1,4 linked glucose monomers having side chains which may include sugars other than glucose. These side chains frequently include xylose, fucose, arabinose, and galactose. Pectins are another type of polysaccharide found in plant cell walls. Pectins are acidic polysaccharides, which are generally comprised primarily of galacturonic acid and rhamnose sugar monomers. Amylose is an additional common plant polysaccharide which is not usually found as a major component of cell walls. It acts primarily as a storage material for glucose, rather than as a structural polymer. However, because amylose is comprised primarily of  $\alpha$ -1,4-linked glucose monomers, it is considered to be a related polymer from a biochemical and physiological perspective.

Plant polysaccharides have many uses. Certain plastics, such as cellulose acetate, and synthetic textiles, such as rayon, are made from cellulose. In addition, some biodegradable plastics and digestible medicine capsules, as well as medical fillers and fiber additives for food, can be made from plant polysaccharides.

30

In foodstuffs, polysaccharides have a profound impact on food quality. Cell walls contribute to crispness in carrots, while degradation of cell walls is required for softening of fruits, such as peaches and tomatoes. In maize, increased amylose is desirable for cattle feed, but not for human consumption, and increased cell wall strength reduces digestibility. In fiber crops, such as timber, cellulose is the primary polymer of interest. Wood density, a fundamental measure of structural timber quality, is essentially a measure of cellulose content. In the paper pulping industry, efficiency is measured in terms of yield of cellulose. Clearly, the ability to increase cellulose content in timber is an important economic goal.

The sugars which make up plant cell wall polysaccharides are produced in the photosynthetic organs of plants. The sugars so produced are commonly converted into sucrose, a disaccharide consisting of glucose and fructose. Sucrose is transported throughout the plant, to wherever sugar monomers are called for. Thus, the photosynthetic organs are often referred to as a source, while tissues requiring large amounts of sugar monomers are referred to as a sink. Actively growing regions of the plant are generally sink tissues, as new cell wall synthesis requires large amounts of sugar monomers.

When the transported sucrose arrives at the sink destination, it must be converted into whichever kind of sugar monomer is required. The sugar monomers which make up plant cell walls are primarily 5- or 6-carbon sugars. Different sugars are generally distinguished by stereospecific orientation of hydroxyl groups. Plants contain a variety of enzymes, such as isomerases or epimerases, which can rapidly change the orientation of these hydroxyls. In addition, there are a number of enzymes which can add or remove a single carbon from a sugar monomer. The result is a single pool of sugar monomers which the plant can freely inter-convert into whichever kind is needed for cell wall synthesis.

Plant polysaccharides are thus biochemically and physiologically inter-related. All polymers compete for the same pool of sugar monomers, and all sugar monomers can be freely interconverted to other types. Degradation of any one polymer will provide building material for any other. Attempts to engineer changes in one polymer may therefore have pleiotropic effects on other polymers.

The rate of cell wall synthesis is dependent on both the availability of sugar monomers to serve as building blocks for the polymers of the wall, and the enzymes which polymerise those building blocks into polymers. Enzymes which are directly responsible for the

synthesis of the major cell wall polymers, such as cellulose, hemicellulose and pectin, may have a profound impact on the rate of cell wall synthesis. Source-sink relations may play an important role in limiting cell wall synthesis, if the availability of substrates becomes limiting. Polymer degrading enzymes may liberate sugar monomers from unnecessary polymers for use  
5 in building new, desired polymers. Enzymes which can isomerise sugars from one form into another can convert the sugars into whichever kind is needed. Each of the different types of cell wall polysaccharides effectively competes for the same pool of sugar monomers, and each represents a potential source of monomers for any of the other polymers.

The final committed steps in cellulose biosynthesis involve a relatively small number  
10 of enzymes. Cellulose synthase (CEL) is believed to function as part of a large, membrane-bound complex which also includes sucrose synthase (SUS: Amor et al., *Proc. Natl. Acad. Sci USA* 92:9353-9357, 1995) and annexin (ANX: Clark and Roux, *Plant Phys.* 109:1133-1139, 1995). This enzyme complex polymerises activated glucose into the cellulose polymer. The glucose is activated by UDP-glucose pyrophosphorylase (UGP), also known as UTP-  
15 glucose-1-phosphate uridylyltransferase. These enzymes are believed to be sufficient for the biosynthesis of cellulose from glucose. Other than these steps, the availability of glucose appears to be the most significant rate-limiting step in cellulose biosynthesis.

Glucose is primarily stored in most plants as amylose. Plants routinely store amylose and degrade it to free up the glucose monomers, as needed. By inhibiting the efficiency of  
20 glucose storage, or by increasing the liberation of glucose from amylose, the availability of glucose monomers for cellulose biosynthesis can be increased. The rate-limiting enzyme in the storage of glucose as amylose is ADP-glucose pyrophosphorylase (AGP), also known as ATP-glucose-1-phosphate adenylyltransferase (Iglesias et al., *J. Biol. Chem.* 268:1081-1086, 1993). Conversely, the enzyme most responsible for liberating glucose from amylose is  
25 amylase (AMA: Kawagoe and Delmer, *Genetic Engineering* 19:63-87, 1997).

These enzymes clearly will be important in the engineering of economically useful changes in cellulose biosynthesis. In addition, there are many other enzymes which may be useful in influencing plant cell wall polysaccharide biosynthesis. Other enzymes likely to be involved in cellulose biosynthesis include 1,4- $\beta$ -cellobiohydrolase,  $\beta$ -glucosidase, calnexin,  
30 cellobiose epimerase, cellobiose phosphorylase, cellulase A, dextranucrase, invertase, phosphodiesterase, phosphoglucomutase, sucrose phosphate synthase, sucrose phosphorylase,

UDP-glucose 4-epimerase and UDP-glucose dehydrogenase. Enzymes believed to be involved in hemicellulose biosynthesis include  $\beta$ -glucanase, arabinan synthase, GDP-fucose pyrophosphorylase, GDP-mannose pyrophosphorylase, 1,3 and 1,4- $\beta$ -glucanases, 1,3 and 1,4- $\beta$ -glucosidases, mannose-6-phosphate isomerase,  $\alpha$ -DP-hexose pyrophosphorylase, xyloglucan endotransglycosylase and xyloglucan synthase. Enzymes likely to be involved in pectin biosynthesis include  $\alpha$ -galactosidase,  $\beta$ -glucuronidase, exopolygalacturonase, glucuronosyl-transferase, pectin methyl-esterase, polygalacturonase and UDP-hexose-1-phosphate uridylyltransferase. Enzymes believed to be involved in amylose biosynthesis include  $\alpha$ -glucosidase, amylopectin 6-glucanohydrolase, amylopectin-branching glycosyltransferase,  $\beta$ -amylase, branching enzyme, inulosucrase, isoamylase, isomaltase, levansucrase, starch phosphorylase and starch synthase. Enzymes likely to be involved in the interconversion of 5-carbon sugars include 2-dehydro-3-deoxy-gluconokinase, aldehyde reductase, arabinose isomerase, D-arabinitol dehydrogenase, D-xylulose reductase, endo-1,4- $\beta$ -xylanase, exo-1,4- $\beta$ -xylanase, L-arabinose isomerase, L-ribulokinase, L-xylulokinase, phospho-ribulokinase, ribose 5-phosphate isomerase, ribulose-phosphate-3-epimerase, ribulose-phosphate-4-epimerase, transaldolase, transketolase, xylose isomerase and xylulokinase. Enzymes likely to be involved in interconversion of 6-carbon sugars include 6-phospho-fructo-1-kinase, 6-phospho-fructo-2-kinase, trehalose phosphate synthase, aldolase, aldose 1-epimerase, D-fructokinase, D-galactokinase, fructose 1,6-diphosphatase, gluconolactonase, glucose 1-phosphatase, glucose 6-phosphatase, glucose 6-phosphate dehydrogenase, glucose-phosphate isomerase, hexokinase, phosphoglucomutase, trehalase, trehalose phosphatase and UDP-galactose dehydrogenase.

While DNA sequences encoding some of the enzymes involved in the biosynthetic pathways of plant cell wall polysaccharides have been isolated for certain species of plants, genes encoding many of the enzymes in a wide range of plant species have not yet been identified. Thus, there remains a need in the art for materials useful in the modification of cell wall polysaccharide content and composition in plants.



### Summary of the Invention

Briefly, the present invention provides polynucleotides isolated from eucalyptus and pine which encode enzymes involved in the synthesis of cell wall polysaccharides. Genetic constructs including such sequences and methods for the use of such constructs are also provided, together with transgenic plants having altered cell wall polysaccharide content and composition.

In one embodiment, the isolated polynucleotides comprise a nucleotide sequence selected from the group consisting of: (a) sequences recited in SEQ ID NOS: 1-29, 57-80, 105, 107, 109-113, 119-129, 139-143 and 149-908; (b) complements of the sequences recited in SEQ ID NOS: 1-29, 57-80, 105, 107, 109-113, 119-129, 139-143 and 149-908; (c) reverse complements of the sequences recited in SEQ ID NOS: 1-29, 57-80, 105, 107, 109-113, 119-129, 139-143 and 149-908; (d) reverse sequences of the sequences recited in SEQ ID NOS: 1-29, 57-80, 105, 107, 109-113, 119-129, 139-143 and 149-908; and (e) sequences having either 40%, 60%, 75% or 90% identical nucleotides, as defined herein, to a sequence of (a) – (d).

In a further aspect, isolated polypeptides encoded by a polynucleotide of the present invention are provided. In one embodiment, such polypeptides comprise an amino acid sequence selected from the group consisting of SEQ ID NOS: 30-56, 81-104, 106, 108, 114-118, 129-138 and 144-148, and variants thereof.

In another aspect, the invention provides genetic constructs comprising a polynucleotide of the present invention, either alone, in combination with one or more of the inventive polynucleotide sequences, or in combination with one or more known polynucleotides, together with transgenic cells comprising such constructs.

In a related aspect, the present invention provides genetic constructs comprising, in the 5'-3' direction, a gene promoter sequence; an open reading frame coding for at least a functional portion of an enzyme encoded by a polynucleotide of the present invention or a variant thereof; and a gene termination sequence. The open reading frame may be orientated in either a sense or antisense direction. Genetic constructs comprising a non-coding region of a gene coding for an enzyme encoded by the above polynucleotides or a nucleotide sequence complementary to a non-coding region, together with a gene promoter sequence and a gene termination sequence, are also provided. Preferably, the gene promoter and termination

sequences are functional in a host plant. Most preferably, the gene promoter and termination sequences are those of the original enzyme genes but others generally used in the art, such as the Cauliflower Mosaic Virus (CMV) promoter, with or without enhancers such as the Kozak sequence or Omega enhancer, and *Agrobacterium tumefaciens* nopal synthase terminator  
5 may be usefully employed in the present invention. Tissue-specific promoters may be employed in order to target expression to one or more desired tissues. In a preferred embodiment, the gene promoter sequence provides for transcription in xylem. The genetic construct may further include a marker for the identification of transformed cells.

In a further aspect, transgenic plant cells comprising the genetic constructs of the  
10 present invention are provided, together with plants comprising such transgenic cells, and fruits, seeds and other products, derivatives, or progeny of such forestry plants. Propagules of the transgenic plants transformed with the inventive polynucleotides are also included in the present invention. As used herein, the word "propagule" means any part of a plant that may be used in reproduction or propagation, sexual or asexual, including cuttings.

15 Plant varieties, particularly registrable plant varieties according to Plant Breeders' Rights, may be excluded from the present invention. A plant need not be considered a "plant variety" simply because it contains stably within its genome a transgene, introduced into a cell of the plant or an ancestor thereof.

In yet another aspect, methods for modulating the polysaccharide content and  
20 composition of an organism, such as a plant, are provided, such methods including stably incorporating into the genome of the plant a genetic construct of the present invention. In a preferred embodiment, the target plant is a woody plant, preferably selected from the group consisting of eucalyptus, pine, acacia, poplar, sweetgum, teak and mahogany species, more preferably from the group consisting of pine and eucalyptus species, and most preferably from  
25 the group consisting of *Eucalyptus grandis* and *Pinus radiata*. In a related aspect, a method for producing a plant having modified cellulose content is provided, the method comprising transforming a plant cell with a genetic construct of the present invention to provide a transgenic cell and cultivating the transgenic cell under conditions conducive to regeneration and mature plant growth.

30 In yet a further aspect, the present invention provides methods for modifying the activity of a polypeptide in a plant, comprising stably incorporating into the genome of the

plant a genetic construct of the present invention. In a preferred embodiment, the target plant is a woody plant, preferably selected from the group consisting of eucalyptus, pine, acacia, poplar, sweetgum, teak and mahogany species, more preferably from the group consisting of pine and eucalyptus species, and most preferably from the group consisting of *Eucalyptus grandis* and *Pinus radiata*.

The above-mentioned and additional features of the present invention and the manner of obtaining them will become apparent, and the invention will be best understood by reference to the following more detailed description. All references disclosed herein are hereby incorporated by reference in their entirety as if each was incorporated individually.

#### Brief Description of the Figures

Fig. 1 illustrates the level of native CEL enzyme activity in positive control mung bean (*V. radiata*) plants.

Fig. 2 illustrates the level of CEL enzyme activity in mammalian 293T cells transfected with *E. grandis* CEL as compared to that in non-transfected 293T cells.

#### Detailed Description

As outlined above, cellulose is formed by polymerization of glucose into a linear, unbranched, polymer comprised of  $\beta$ -1,4 linked glucose monomers (Kawagoe and Delmer, *Genetic Engineering*, 19:63-87, 1997). Cellulose is the most important plant cell wall polysaccharide from both a structural, as well as industrial, perspective. Other polysaccharides are essential for healthy cell walls, as well as for many alternative industrial uses.

Glucose monomers are most commonly stored in the plant in the form of amylose by the action of several enzymes, with the rate limiting step for storage being catalysed by AGP (Iglesias et al., *J. Biol. Chem.* 268:1081-1086). Glucose monomers are freed from this storage polymer by the action of the enzyme AMA. The free monomers are activated by the action of the enzyme UGP, and polymerised into cellulose macro-crystalline structures by the action of the cellulose synthase enzyme complex. Pure CEL enzyme has been shown to form  $\beta$ -1,4 glucose linkages *in vitro*, but has not been shown to be sufficient for polymerization of the

large polymers which are fundamental to the structure of plant cell walls. The holoenzyme complex appears to be necessary for this latter function. The holoenzyme is believed to be comprised of the CEL enzyme in combination with the SUS enzyme and ANX, the whole complex being integrated into the plasma membrane and forming a "rosette" structure as seen  
5 in electron micrographs of plant cell membranes (Arioli et al., *Science* 279:717-720, 1998).

Because cellulose synthesis can represent such a large sink for sugar monomers in the cell, changes in the rate of cellulose synthesis can have a profound influence on the synthesis of other plant polysaccharides. Conversely, changes in the rates of synthesis of other plant polysaccharides can have a profound influence on the pool of sugars available for synthesis of  
10 cellulose. Hence, changes in the synthesis of any single polymer may affect both the content and composition of plant cell wall polysaccharides, and polysaccharides in general.

Quantitative and qualitative modifications in plant polysaccharide content are known to be induced by external factors such as light stimulation, low calcium levels, and mechanical stress. Synthesis of cell wall polysaccharides can also be induced by infection  
15 with pathogens.

Using the methods and materials of the present invention, the polysaccharide content of a plant may be increased or reduced, by incorporating additional copies of genes encoding enzymes involved in the synthesis of cell wall polysaccharides into the genome of the target plant. Similarly, an increase or decrease in polysaccharide content may be obtained by  
20 transforming the target plant with antisense copies of such genes. In addition, the number of copies of genes encoding for different enzymes in the biosynthetic pathway of cell wall polysaccharides can be manipulated to modify the relative amount of each monosaccharide synthesized, thereby leading to the formation of cell walls having altered composition. The alteration of polysaccharide composition would be advantageous, for example, in tree  
25 processing for paper.

The polynucleotides of the present invention were isolated from forestry plant sources, namely from *Eucalyptus grandis* and *Pinus radiata*, but they may alternatively be synthesized using conventional synthesis techniques. Specifically, isolated polynucleotides of the present invention include polynucleotides comprising a sequence selected from the group consisting  
30 of sequences identified as SEQ ID NOS: 1-29, 57-80, 105, 107, 109-113, 119-129, 139-143 and 149-908; complements of the sequences identified as SEQ ID NOS: 1-29, 57-80, 105,

107, 109-113, 119-129, 139-143 and 149-908; reverse complements of the sequences identified as SEQ ID NOS: 1-29, 57-80, 105, 107, 109-113, 119-129, 139-143 and 149-908; at least a specified number of contiguous residues (x-mers) of any of the above-mentioned polynucleotides; extended sequences corresponding to any of the above polynucleotides; antisense sequences corresponding to any of the above polynucleotides; and variants of any of the above polynucleotides, as that term is described in this specification.

In another embodiment, the present invention provides isolated polypeptides encoded by the DNA sequences of SEQ ID NOS: 1-29, 57-80, 105, 107, 109-113, 119-129, 139-143 and 149-908;. The predicted amino acid sequences encoded by SEQ ID NOS: 1-22, 24-28, 57-80, 105, 107, 109-113 and 119-143, based on the best available information at the time of filing this application, are provided in SEQ ID NOS: 30-56, 81-104, 106, 108, 114-118, 129-138 and 144-148, respectively. The present invention also encompasses polynucleotides that differ from the disclosed sequences but which, due to the degeneracy of the genetic code, encode a polypeptide which is the same as that encoded by a polypeptide of the present invention. Such polynucleotides are said to be "degeneratively equivalent" to a polynucleotide sequence disclosed herein.

The polynucleotides and polypeptides of the present invention were putatively identified by DNA and polypeptide similarity searches. In the attached Sequence Listing SEQ ID NOS: 1-29, 57-80, 105, 107, 109-113, 119-129, 139-143 and 149-908 are polynucleotide sequences, and SEQ ID NOS: 30-56, 81-104, 106, 108, 114-118, 129-138 and 144-148 are polypeptide sequences. The polynucleotides and polypeptides of the present invention, have demonstrated similarity to enzymes that are known to be involved in the synthesis of cell wall polysaccharides. The putative identity of each of the inventive polynucleotides is shown below in Table 1.

TABLE 1

DNA SEQ ID NO:	PROTEIN SEQ ID NO:	IDENTITY
1	30	AGP
2	31	AGP
3	32	AGP
4	33	AMA

DNA SEQ ID NO:	PROTEIN SEQ ID NO:	IDENTITY
5	34	AMA
6	35	AMA
7	36	CEL
8	37	CEL
9	38	CEL
10	39	CEL
11	40	CEL
12	41	CEL
13	42	CEL
14	43	CEL
15	44	SUS
16	45	SUS
17	46	SUS
18	47	SUS
19	48	SUS
20	49	UGP
21	50	UGP
22	51	UGP
23	-	UGP
24	52	ANX
25	53	ANX
26	54	ANX
27	55	ANX
28	56	ANX
29	-	ANX
57	81	AMA
58	82	AMA
59	83	AGP
60	84	AGP
61	85	AGP
62	86	AGP
63	87	AGP
64	88	AGP
65	89	AGP
66	90	CEL
67	91	CEL
68	92	CEL
69	93	CEL
70	94	CEL
71	95	SUS
72	96	SUS
73	97	SUS
74	98	SUS
75	99	SUS
76	100	SUS
77	101	SUS
78	102	SUS
79	103	UGP
80	104	UGP

DNA SEQ ID NO:	PROTEIN SEQ ID NO:	IDENTITY
105	106	SUS
107	108	CEL
109	114	ANX
110	115	ANX
111	116	ANX
112	117	ANX
113	118	ANX
119	129	CEL
120	130	CEL
121	131	CEL
122	132	CEL
123	133	CEL
124	134	CEL
125	135	CEL
126	136	CEL
127	137	CEL
128	138	CEL
135	144	SUS
140	145	$\alpha$ -amylase
141	146	CEL
142	147	AGP (3' end of SEQ ID NO: 62)
143	148	SUS (3' of SEQ ID NO: 74)
149-185	-	1,3- $\beta$ -D-Glucanase
186	-	1,4- $\beta$ -Cellobiohydrolase
187-196	-	$\alpha,\alpha$ -trehalose phosphate synthase
197-204	-	$\alpha$ -glucosidase
205-250	-	aldolase
251	-	Amylopectin 6-glucanohydrolase
252-262	-	$\beta$ -amylase
263	-	$\beta$ -glucosidase
264-272	-	Branching enzyme
273-318	-	D-fructokinase
319-354	-	D-xylulose reductase
355-365	-	Endo-1,3-1,4- $\beta$ -glucanase
366-371	-	Glucan exo-1,3- $\beta$ -glucosidase
372-377	-	Glucose 6-phosphate dehydrogenase
378-381	-	Glucose phosphate isomerase
382-389	-	Isoamylase
390-393	-	L-ribulokinase
394-398	-	Mannitol-1-phosphate 5-dehydrogenase
399-478	-	Pectin methyl-esterase
479-506	-	Phosphoglucomutase
507-508	-	Phospho-ribulokinase
509-521	-	Ribulose-phosphate-3-epimerase
522-530	-	Starch phosphorylase
531-551	-	Sucrose phosphate synthase
552-555	-	SUS
556-586	-	Transketolase
587-591	-	Trehalase

DNA SEQ ID NO:	PROTEIN SEQ ID NO:	IDENTITY
592-620	-	UDP-glucose 4-epimerase
621-902	-	Xyloglucan endotransglycosylase
903-908	-	Xylose isomerase

The term "polynucleotide(s)," as used herein, means a single or double-stranded polymer of deoxyribonucleotide or ribonucleotide bases and includes DNA and corresponding RNA molecules, including HnRNA and mRNA molecules, both sense and anti-sense strands, and comprehends cDNA, genomic DNA and recombinant DNA, as well as wholly or partially synthesized polynucleotides. An HnRNA molecule contains introns and corresponds to a DNA molecule in a generally one-to-one manner. An mRNA molecule corresponds to an HnRNA and DNA molecule from which the introns have been excised. A polynucleotide may consist of an entire gene, or any portion thereof. Operable anti-sense polynucleotides may comprise a fragment of the corresponding polynucleotide, and the definition of "polynucleotide" therefore includes all such operable anti-sense fragments.

The term "polypeptide", as used herein, encompasses amino acid chains of any length including full length proteins, wherein amino acid residues are linked by covalent peptide bonds. Polypeptides of the present invention may be naturally purified products, or may be produced partially or wholly using recombinant techniques.

The definition of the terms "complement", "reverse complement" and "reverse sequence", as used herein, is best illustrated by the following example. For the sequence 5' AGGACC 3', the complement, reverse complement and reverse sequence are as follows:

complement	3' TCCTGG 5'
reverse complement	3' GGTCCT 5'
reverse sequence	5' CCAGGA 3'.

As used herein, the term "variant" covers any sequence which has at least about 40%, more preferably at least about 60%, more preferably yet at least about 75% and most preferably at least about 90% identical residues (either nucleotides or amino acids) to a sequence of the present invention. The percentage of identical residues is determined by aligning the two sequences to be compared, determining the number of identical residues in



the aligned portion, dividing that number by the total length of the inventive, or queried, sequence and multiplying the result by 100.

Polynucleotide or polypeptide sequences may be aligned, and percentage of identical nucleotides in a specified region may be determined against another polynucleotide, using computer algorithms that are publicly available. Two exemplary algorithms for aligning and identifying the similarity of polynucleotide sequences are the BLASTN and FASTA algorithms. The similarity of polypeptide sequences may be examined using the BLASTP algorithm. Both the BLASTN and BLASTP software are available on the NCBI anonymous FTP server (<ftp://ncbi.nlm.nih.gov>) under /blast/executables/. The BLASTN algorithm Version 2.0.6 [Sept-16-1998], set to the default parameters described in the documentation and distributed with the algorithm, is preferred for use in the determination of variants according to the present invention. The use of the BLAST family of algorithms, including BLASTN and BLASTP, is described at NCBI's Internet website at the URL <http://www.ncbi.nlm.nih.gov/BLAST/newblast.html> and in the publication of Altschul, Stephen F, et al., "Gapped BLAST and PSI-BLAST: a new generation of protein database search programs," *Nucleic Acids Res.* 25:3389-3402, 1997. The computer algorithm FASTA is available on the Internet at the ftp site <ftp://ftp.virginia.edu/pub/fasta/>. Version 2.04, [February 1996], set to the default parameters described in the documentation and distributed with the algorithm, is preferred for use in the determination of variants according to the present invention. The use of the FASTA algorithm is described in Pearson WR and Lipman DJ, "Improved Tools for Biological Sequence Analysis," *Proc. Natl. Acad. Sci. USA* 85:2444-2448, 1988; and Pearson WR, "Rapid and Sensitive Sequence Comparison with FASTP and FASTA," *Methods in Enzymol.* 183:63-98, 1990.

The following running parameters are preferred for determination of alignments and identities using BLASTN that contribute to the E values and percentage identity of polynucleotides of the present invention: Unix running command: `blastall -p blastn -d embldb -e 10 -G 0 -E 0 -r 1 -v 30 -b 30 -i queryseq -o results`; and the parameters are: -p Program Name [String]; -d Database [String]; -e Expectation value (E) [Real]; -G Cost to open a gap (zero invokes default behavior) [Integer]; -E Cost to extend a gap (zero invokes default behavior) [Integer]; -r Reward for a nucleotide match (blastn only) [Integer]; -v Number of

one-line descriptions (V) [Integer]; -b Number of alignments to show (B) [Integer]; -i Query File [File In]; -o BLAST report Output File [File Out] Optional.

The following running parameters are preferred for determination of alignments and identities using BLASTP that contribute to the E values and percentage identity of polypeptide sequences: For BLASTP the following running parameters are preferred: blastall -p blastp -d swissprot -e 10 -G 0 -E 0 -v 30 -b 30 -i queryseq -o results; and parameters are: -p Program Name [String]; -d Database [String]; -e Expectation value (E) [Real]; -G Cost to open a gap (zero invokes default behavior) [Integer]; -E Cost to extend a gap (zero invokes default behavior) [Integer]; -v Number of one-line descriptions (v) [Integer]; -b Number of alignments to show (b) [Integer]; -I Query File [File In]; -o BLAST report Output File [File Out] Optional.

The "hits" to one or more database sequences by a queried sequence produced by BLASTN, BLASTP, FASTA, or a similar algorithm, align and identify similar portions of sequences. The hits are arranged in order of the degree of similarity and the length of sequence overlap. Hits to a database sequence generally represent an overlap over only a fraction of the sequence length of the queried sequence.

The BLASTN and FASTA algorithms also produce "Expect" values for alignments. The Expect value (E) indicates the number of hits one can "expect" to see over a certain number of contiguous sequences by chance when searching a database of a certain size. The Expect value is used as a significance threshold for determining whether the hit to a database, such as the preferred EMBL database, indicates true similarity. For example, an E value of 0.1 assigned to a hit is interpreted as meaning that in a database of the size of the EMBL database, one might expect to see 0.1 matches over the aligned portion of the sequence with a similar score simply by chance. By this criterion, the aligned and matched portions of the sequences then have a probability of 90% of being the same. For sequences having an E value of 0.01 or less over aligned and matched portions, the probability of finding a match by chance in the EMBL database is 1% or less using the BLASTN or FASTA algorithm.

According to one embodiment, "variant" polynucleotides, with reference to each of the polynucleotides of the present invention, preferably comprise sequences having the same number or fewer nucleic acids than each of the polynucleotides of the present invention and producing an E value of 0.01 or less when compared to the polynucleotide of the present

invention. That is, a variant polynucleotide is any sequence that has at least a 99% probability of being the same as the polynucleotide of the present invention, measured as having an E value of 0.01 or less using the BLASTN or FASTA algorithms set at the default parameters. According to a preferred embodiment, a variant polynucleotide is a sequence having the same  
5 number or fewer nucleic acids than a polynucleotide of the present invention that has at least a 99% probability of being the same as the polynucleotide of the present invention, measured as having an E value of 0.01 or less using the BLASTN or FASTA algorithms set at the default parameters.

Alternatively, variant polynucleotide hybridize to the polynucleotide of the present  
10 invention under stringent conditions. As used herein, "stringent conditions" refers to prewashing in a solution of 6X SSC, 0.2% SDS; hybridizing at 65°C, 6X SSC, 0.2% SDS overnight; followed by two washes of 30 minutes each in 1X SSC, 0.1% SDS at 65°C and two washes of 30 minutes each in 0.2X SSC, 0.1% SDS at 65°C.

The present invention also encompasses polynucleotides that differ from the disclosed  
15 sequences but that, as a consequence of the discrepancy of the genetic code, encode a polypeptide having similar enzymatic activity as a polypeptide encoded by a polynucleotide of the present invention. Thus, polynucleotides comprising sequences that differ from the polynucleotide sequences recited in SEQ ID NOS: 1-29, 57-80, 105, 107, 109-113, 119-129, 139-143 and 149-908, or complements, reverse sequences, or reverse complements of those  
20 sequences as a result of conservative substitutions are contemplated by and encompassed within the present invention. Additionally, polynucleotides comprising sequences that differ from the polynucleotide sequences recited in SEQ ID NOS: 1-29, 57-80, 105, 107, 109-113, 119-129, 139-143 and 149-908, or complements, reverse complements, or reverse sequences as a result of deletions and/or insertions totaling less than 10% of the total sequence length are  
25 also contemplated by and encompassed within the present invention. Similarly, polypeptides comprising sequences that differ from the polypeptide sequences recited in SEQ ID NOS: 30-56, 81-104, 106, 108, 114-118, 129-138 and 144-148 as a result of amino acid substitutions, insertions, and/or deletions totaling less than 10% of the total sequence length are contemplated by an encompassed within the present invention, provided the variant  
30 polypeptide has activity in a cell wall polysaccharide synthesis pathway.

5 Variants of the polypeptide sequences recited in SEQ ID NOS: 30-56, 81-104, 106, 108, 114-118, 129-138 and 144-148, wherein the variant has an activity level that is different to that of the recited polypeptide are also encompassed by the present invention. In specific embodiments, variants of the inventive sucrose synthase (SUS) polypeptides are provided wherein the N-terminal serine phosphorylation site has been replaced by an acidic amino acid (such as Asp or Glu) by, for example, site directed mutagenesis. Nakai et al. have demonstrated that SUS polypeptides mutated in this manner possess increased activity compared to wild-type SUS (Nakai et al., *Plant Cell Physiol.* 39:1337-1341, 1998). Polynucleotides encoding such variants of the inventive SUS polypeptides may therefore be employed in transgenic plants to increase cellulose production.

10 The polynucleotides of the present invention may be isolated from various libraries, or may be synthesized using techniques that are well known in the art. The polynucleotides may be synthesized, for example, using automated oligonucleotide synthesizers (e.g., Beckman Oligo 1000M DNA Synthesizer) to obtain polynucleotide segments of up to 50 or more nucleic acids. A plurality of such polynucleotide segments may then be ligated using standard DNA manipulation techniques that are well known in the art of molecular biology. One conventional and exemplary polynucleotide synthesis technique involves synthesis of a single stranded polynucleotide segment having, for example, 80 nucleic acids, and hybridizing that segment to a synthesized complementary 85 nucleic acid segment to produce a 5 nucleotide overhang. The next segment may then be synthesized in a similar fashion, with a 5 nucleotide overhang on the opposite strand. The "sticky" ends ensure proper ligation when the two portions are hybridized. In this way, a complete polynucleotide of the present invention may be synthesized entirely *in vitro*.

25 Some of the polynucleotides identified as SEQ ID NOS: 1-29, 57-80, 105, 107, 109-113, 119-129, 139-143 and 149-908 are referred to as "partial" sequences, in that they do not represent the full coding portion of a gene encoding a naturally occurring polypeptide. The partial polynucleotide sequences disclosed herein may be employed to obtain the corresponding full length genes for various species and organisms by, for example, screening DNA expression libraries using hybridization probes based on the polynucleotides of the present invention, or using PCR amplification with primers based upon the polynucleotides of the present invention. In this way one can, using methods well known in the art, extend a

polynucleotide of the present invention upstream and downstream of the corresponding mRNA, as well as identify the corresponding genomic DNA, including the promoter and enhancer regions, of the complete gene. The present invention thus comprehends isolated polynucleotides comprising a sequence identified in SEQ ID NOS: 1-29, 57-80, 105, 107, 109-113, 119-129, 139-143 and 149-908, or a variant of one of the specified sequences, that encode a functional polypeptide, including full length genes. Such extended polynucleotides may have a length of from about 50 to about 4,000 nucleic acids or base pairs, and preferably have a length of less than about 4,000 nucleic acids or base pairs, more preferably yet a length of less than about 3,000 nucleic acids or base pairs, more preferably yet a length of less than about 2,000 nucleic acids or base pairs. Under some circumstances, extended polynucleotides of the present invention may have a length of less than about 1,800 nucleic acids or base pairs, preferably less than about 1,600 nucleic acids or base pairs, more preferably less than about 1,400 nucleic acids or base pairs, more preferably yet less than about 1,200 nucleic acids or base pairs, and most preferably less than about 1,000 nucleic acids or base pairs.

Polynucleotides of the present invention also comprehend polynucleotides comprising at least a specified number of contiguous residues ( $x$ -mers) of any of the polynucleotides identified as SEQ ID NOS: 1-29, 57-80, 105, 107, 109-113, 119-129, 139-143 and 149-908, complements, reverse sequences, and reverse complements of such sequences, and their variants. Similarly, polypeptides of the present invention comprehend polypeptides comprising at least a specified number of contiguous residues ( $x$ -mers) of any of the polypeptides identified as SEQ ID NOS: 30-56, 81-104, 106, 108, 114-118, 129-138 and 144-148, and their variants. As used herein, the term " $x$ -mer," with reference to a specific value of " $x$ ," refers to a sequence comprising at least a specified number (" $x$ ") of contiguous residues of any of the polynucleotides identified as SEQ ID NO: 1-29, 57-80, 105, 107, 109-113, 119-129, 139-143 and 149-908, or the polypeptides identified as SEQ ID NOS: 30-56, 81-104, 106, 108, 114-118, 129-138 and 144-148. According to preferred embodiments, the value of  $x$  is preferably at least 20; more preferably, at least 40; more preferably yet, at least 60; and most preferably, at least 80. Thus, polynucleotides and polypeptides of the present invention comprise a 20-mer, a 40-mer, a 60-mer, an 80-mer, a 100-mer, a 120-mer, a 150-mer, a 180-mer, a 220-mer, a 250-mer, or a 300-mer, 400-mer,

500-mer or 600-mer of a polynucleotide or polypeptide identified as SEQ ID NOS: 1-908, and variants thereof.

Polynucleotide probes and primers complementary to and/or corresponding to SEQ ID NOS: 1-29, 57-80, 105, 107, 109-113, 119-129, 139-143 and 149-908, and variants of those  
5 sequences, are also comprehended by the present invention. Such oligonucleotide probes and primers are substantially complementary to the polynucleotide of interest. As used herein, the term "oligonucleotide" refers to a relatively short segment of a polynucleotide sequence, generally comprising between 6 and 60 nucleotides, and comprehends both probes for use in hybridization assays and primers for use in the amplification of DNA by polymerase chain  
10 reaction.

An oligonucleotide probe or primer is described as "corresponding to" a polynucleotide of the present invention, including one of the sequences set out as SEQ ID NOS: 1-29, 57-80, 105, 107, 109-113, 119-129, 139-143 and 149-908, or a variant, if the oligonucleotide probe or primer, or its complement, is contained within one of the sequences  
15 set out as SEQ ID NOS: 1-29, 57-80, 105, 107, 109-113, 119-129, 139-143 and 149-908, or a variant of one of the specified sequences.

Two single stranded sequences are said to be substantially complementary when the nucleotides of one strand, optimally aligned and compared, with the appropriate nucleotide insertions and/or deletions, pair with at least 80%, preferably at least 90% to 95%, and more  
20 preferably at least 98% to 100%, of the nucleotides of the other strand. Alternatively, substantial complementarity exists when a first DNA strand will selectively hybridize to a second DNA strand under stringent hybridization conditions. Stringent hybridization conditions for determining complementarity include salt conditions of less than about 1 M, more usually less than about 500 mM, and preferably less than about 200 mM. Hybridization  
25 temperatures can be as low as 5°C, but are generally greater than about 22°C, more preferably greater than about 30°C, and most preferably greater than about 37°C. Longer DNA fragments may require higher hybridization temperatures for specific hybridization. Since the stringency of hybridization may be affected by other factors such as probe composition, presence of organic solvents and extent of base mismatching, the combination of parameters  
30 is more important than the absolute measure of any one alone. The DNA from plants or

samples or products containing plant material can be either genomic DNA or DNA derived by preparing cDNA from the RNA present in the sample.

In addition to DNA-DNA hybridization, DNA-RNA or RNA-RNA hybridization assays are also possible. In the first case, the mRNA from expressed genes would then be  
5 detected instead of genomic DNA or cDNA derived from mRNA of the sample. In the second case, RNA probes could be used. In addition, artificial analogs of DNA hybridizing specifically to target sequences could also be used.

In specific embodiments, the oligonucleotide probes and/or primers comprise at least about 6 contiguous residues, more preferably at least about 10 contiguous residues, and most  
10 preferably at least about 20 contiguous residues complementary to a polynucleotide sequence of the present invention. Probes and primers of the present invention may be from about 8 to 100 base pairs in length or, preferably from about 10 to 50 base pairs in length or, more preferably from about 15 to 40 base pairs in length. The probes can be easily selected using procedures well known in the art, taking into account DNA-DNA hybridization stringencies,  
15 annealing and melting temperatures, and potential for formation of loops and other factors, which are well known in the art. Tools and software suitable for designing probes, and especially suitable for designing PCR primers, are available on the Internet, for example, at URL <http://www.horizonpress.com/pcr/>. Preferred techniques for designing PCR primers are also disclosed in Dieffenbach CW and Dykster GS, *PCR primer: a laboratory manual*,  
20 CSHL Press: Cold Spring Harbor, NY, 1995.

A plurality of oligonucleotide probes or primers corresponding to a polynucleotide of the present invention may be provided in a kit form. Such kits generally comprise multiple DNA or oligonucleotide probes, each probe being specific for a polynucleotide sequence. Kits of the present invention may comprise one or more probes or primers corresponding to a  
25 polynucleotide of the present invention, including a polynucleotide sequence identified in SEQ ID NOS: 1-29, 57-80, 105, 107, 109-113, 119-129, 139-143 and 149-908.

In one embodiment, the present invention provides genetic that include an open reading frame coding for at least a functional portion of a polypeptide encoded by a polynucleotide of the present invention or a variant thereof. As used herein, the "functional  
30 portion" of a polypeptide is that portion which contains the active site essential for affecting the metabolic step, *i.e.*, the portion of the molecule that is capable of binding one or more

reactants or is capable of improving or regulating the rate of reaction. The functional portion can be determined by targeted mutagenesis and screening of modified protein products with protocols well known in the art. Normally, the functional portion is 10-20 amino acids, but can be shorter or longer. The active site may be made up of separate portions present on one or more polypeptide chains and will generally exhibit high substrate specificity. The term "polypeptide encoded by a polynucleotide" as used herein, includes polypeptides encoded by a nucleotide sequence which includes the partial isolated DNA sequences of the present invention.

The open reading frame may be inserted in the genetic construct in a sense or antisense orientation, such that transformation of a target plant with the genetic construct will produce a change in the amount or structure of the polypeptide compared to the wild-type plant. Transformation with a genetic construct comprising an open reading frame in a sense orientation will generally result in modified expression of the selected gene, while transformation with a genetic construct comprising an open reading frame in an antisense orientation also generally results in modified expression of the selected gene. A population of plants transformed with a genetic construct comprising an open reading frame of the present invention in either a sense or antisense orientation may be screened for increased or reduced expression of the gene in question using techniques well known to those of skill in the art, and plants having the desired phenotypes may thus be isolated.

Alternatively, expression of a gene involved in the biosynthesis of polysaccharides may be inhibited by inserting a portion of an open reading frame of the present invention, in either sense or antisense orientation, in the genetic construct. Such portions need not be full-length but preferably comprise at least 25 and more preferably at least 50 residues of a polynucleotide of the present invention. A much longer portion or even the full length polynucleotide corresponding to the complete open reading frame may be employed. The portion of the open reading frame does not need to be precisely the same as the endogenous sequence, provided that there is sufficient sequence similarity to achieve inhibition of the target gene. Thus a sequence derived from one species may be used to inhibit expression of a gene in a different species.

In a second embodiment, the inventive genetic constructs comprise a polynucleotide including a non-coding region of a gene coding for a polypeptide encoded by a polynucleotide



of the present invention, or a polynucleotide sequence complementary to such a non-coding region. Examples of non-coding regions which may be usefully employed in such constructs include introns and 5'-non-coding leader sequences. Transformation of a target plant with such a genetic construct may lead to a reduction in the amount of polysaccharide synthesized by the plant by the process of co-suppression, in a manner similar to that discussed, for example, by Napoli et al. (*Plant Cell* 2:279-290, 1990) and de Carvalho Niebel et al. (*Plant Cell* 7:347-358, 1995).

Alternatively, regulation of polysaccharide synthesis can be achieved by inserting appropriate sequences or subsequences (e.g., DNA or RNA) in ribozyme constructs (McIntyre CL, Manners JM, *Transgenic Res.* 5(4):257-262, 1996). Ribozymes are synthetic RNA molecules that comprise a hybridizing region complementary to two regions, each of which comprises at least 5 contiguous nucleotides in a mRNA molecule encoded by one of the inventive polynucleotides. Ribozymes possess highly specific endonuclease activity, which autocatalytically cleaves the mRNA.

The genetic constructs of the present invention further comprise a gene promoter sequence and a gene termination sequence, operably linked to the DNA sequence to be transcribed, which control expression of the gene. The gene promoter sequence is generally positioned at the 5' end of the DNA sequence to be transcribed, and is employed to initiate transcription of the DNA sequence. Gene promoter sequences are generally found in the 5' non-coding region of a gene but they may exist downstream of the open reading frame, in introns (Luehrsen KR, *Mol. Gen. Genet.* 225:81-93, 1991) or in the coding region, as for example in a plant defence gene (Douglas et al., *EMBO J.* 10:1767-1775, 1991). When the construct includes an open reading frame in a sense orientation, the gene promoter sequence also initiates translation of the open reading frame. For DNA constructs comprising either an open reading frame in an antisense orientation or a non-coding region, the gene promoter sequence consists only of a transcription initiation site having a RNA polymerase binding site.

A variety of gene promoter sequences which may be usefully employed in the DNA constructs of the present invention are well known in the art. The gene promoter sequence, and also the gene termination sequence, may be endogenous to the target plant host or may be exogenous, provided the promoter is functional in the target host. For example, the promoter and termination sequences may be from other plant species, plant viruses, bacterial plasmids,

and the like. Preferably, gene promoter and termination sequences are from the inventive sequences themselves.

Factors influencing the choice of promoter include the desired tissue specificity of the construct, and the timing of transcription and translation. For example, constitutive  
5 promoters, such as the 35S Cauliflower Mosaic Virus (CaMV 35S) promoter, will affect the activity of the enzyme in all parts of the plant. Use of a tissue specific promoter will result in production of the desired sense or antisense RNA only in the tissue of interest. With genetic constructs employing inducible gene promoter sequences, the rate of RNA polymerase binding and initiation can be modulated by external stimuli, such as light, heat, anaerobic  
10 stress, alteration in nutrient conditions and the like. Temporally regulated promoters can be employed to effect modulation of the rate of RNA polymerase binding and initiation at a specific time during development of a transformed cell. Preferably, the original promoters from the enzyme gene in question, or promoters from a specific tissue-targeted gene in the organism to be transformed, such as eucalyptus or pine are used. Other examples of gene  
15 promoters which may be usefully employed in the present invention include mannopine synthase (mas), octopine synthase (ocs), and those reviewed by Chua et al. (*Science* 244:174-181, 1989).

The gene termination sequence, which is located 3' to the DNA sequence to be transcribed, may come from the same gene as the gene promoter sequence or may be from a  
20 different gene. Many gene termination sequences known in the art may be usefully employed in the present invention, such as the 3' end of the *Agrobacterium tumefaciens* nopaline synthase gene. However, preferred gene terminator sequences are those from the original enzyme gene or from the target species to be transformed.

The genetic constructs of the present invention may also contain a selection marker  
25 that is effective in plant cells, to allow for the detection of transformed cells containing the inventive construct. Such markers, which are well known in the art, typically confer resistance to one or more toxins. One example of such a marker is the NPTII gene whose expression results in resistance to kanamycin or hygromycin, antibiotics which are usually toxic to plant cells at a moderate concentration (Rogers et al., in Weissbach, A and H, eds.,  
30 *Methods for Plant Molecular Biology*, Academic Press Inc.: San Diego, CA, 1988). Transformed cells can thus be identified by their ability to grow in media containing the

antibiotic in question. Alternatively, the presence of the desired construct in transformed cells can be determined by means of other techniques well known in the art, such as Southern and Western blots.

5 A transcription initiation site is additionally included in the genetic construct when the sequence to be transcribed lacks such a site.

Techniques for operatively linking the components of the inventive genetic constructs are well known in the art and include the use of synthetic linkers containing one or more restriction endonuclease sites as described, for example, by Sambrook et al. (*Molecular cloning: a laboratory manual*, CSHL Press: Cold Spring Harbor, NY, 1989). The genetic  
10 construct of the present invention may be linked to a vector having at least one replication system, for example *E. coli*, whereby after each manipulation, the resulting construct can be cloned and sequenced and the correctness of the manipulation determined.

The genetic constructs of the present invention may be used to transform a variety of plants, both monocotyledonous (e.g., grasses, corn, grains, oat, wheat and barley),  
15 dicotyledonous (e.g., *Arabidopsis*, tobacco, legumes, alfalfa, oaks, eucalyptus, maple), and Gymnosperms (e.g., Scots pine; see Aronen, *Finnish Forest Res. Papers*, Vol. 595, 1996), white spruce (Ellis et al., *Biotechnology* 11:84-89, 1993), and larch (Huang et al., *In Vitro Cell* 27:201-207, 1991). In a preferred embodiment, the inventive genetic constructs are employed to transform woody plants, herein defined as a tree or shrub whose stem lives for a  
20 number of years and increases in diameter each year by the addition of woody tissue. Preferably the target plant is selected from the group consisting of eucalyptus and pine species, most preferably from the group consisting of *Eucalyptus grandis* and *Pinus radiata*. Other species which may be usefully transformed with the DNA constructs of the present invention include, but are not limited to: pines such as *Pinus banksiana*, *Pinus brutia*, *Pinus caribaea*, *Pinus clausa*, *Pinus contorta*, *Pinus coulteri*, *Pinus echinata*, *Pinus eldarica*, *Pinus ellioti*, *Pinus jeffreyi*, *Pinus lambertiana*, *Pinus monticola*, *Pinus nigra*, *Pinus palustris*,  
25 *Pinus pinaster*, *Pinus ponderosa*, *Pinus resinosa*, *Pinus rigida*, *Pinus serotina*, *Pinus strobus*, *Pinus sylvestris*, *Pinus taeda*, *Pinus virginiana*; other gymnosperms, such as *Abies amabilis*, *Abies balsamea*, *Abies concolor*, *Abies grandis*, *Abies lasiocarpa*, *Abies magnifica*, *Abies procera*, *Chamaecyparis lawsoniana*, *Chamaecyparis nootkatensis*, *Chamaecyparis thyoides*,  
30 *Huniperus virginiana*, *Larix decidua*, *Larix laricina*, *Larix leptolepis*, *Larix occidentalis*,

*Larix siberica*, *Libocedrus decurrens*, *Picea abies*, *Picea engelmanni*, *Picea glauca*, *Picea mariana*, *Picea pungens*, *Picea rubens*, *Picea sitchensis*, *Pseudotsuga menziesii*, *Sequoia gigantea*, *Sequoia sempervirens*, *Taxodium distichum*, *Tsuga canadensis*, *Tsuga heterophylla*, *Tsuga mertensiana*, *Thuja occidentalis*, *Thuja plicata*; and Eucalypts, such as *Eucalyptus* 5 *alba*, *Eucalyptus bancroftii*, *Eucalyptus botyroides*, *Eucalyptus bridgesiana*, *Eucalyptus calophylla*, *Eucalyptus camaldulensis*, *Eucalyptus citriodora*, *Eucalyptus cladocalyx*, *Eucalyptus coccifera*, *Eucalyptus curtisii*, *Eucalyptus dalrympleana*, *Eucalyptus deglupta*, *Eucalyptus delagatensis*, *Eucalyptus diversicolor*, *Eucalyptus dunnii*, *Eucalyptus ficifolia*, *Eucalyptus globulus*, *Eucalyptus gomphocephala*, *Eucalyptus gunnii*, *Eucalyptus henryi*, 10 *Eucalyptus laevopinea*, *Eucalyptus macarthurii*, *Eucalyptus macrorhyncha*, *Eucalyptus maculata*, *Eucalyptus marginata*, *Eucalyptus megacarpa*, *Eucalyptus melliodora*, *Eucalyptus nicholii*, *Eucalyptus nitens*, *Eucalyptus nova-anglica*, *Eucalyptus obliqua*, *Eucalyptus obtusiflora*, *Eucalyptus oreades*, *Eucalyptus pauciflora*, *Eucalyptus polybractea*, *Eucalyptus regnans*, *Eucalyptus resinifera*, *Eucalyptus robusta*, *Eucalyptus rudis*, *Eucalyptus saligna*, 15 *Eucalyptus sideroxylon*, *Eucalyptus stuartiana*, *Eucalyptus tereticornis*, *Eucalyptus torelliana*, *Eucalyptus urnigera*, *Eucalyptus urophylla*, *Eucalyptus viminalis*, *Eucalyptus viridis*, *Eucalyptus wandoo* and *Eucalyptus youmanni*; together with hybrids of the above species.

As discussed above, transformation of a plant with a genetic construct of the present 20 invention will result in a modification in polysaccharide synthesis in the plant. For example, an increase in the production of cellulose in a plant may be obtained by introducing a genetic construct comprising an open reading frame encoding the enzyme CEL in a sense orientation. Similarly, transformation of a plant with a genetic construct comprising either an open reading frame encoding CEL in an antisense orientation or a non-coding (untranslated) region 25 of a CEL gene will lead to a reduction in the cellulose content of the transformed plant.

Techniques for stably incorporating genetic constructs into the genome of target plants are well known in the art and include *Agrobacterium tumefaciens* mediated introduction, electroporation, protoplast fusion, injection into reproductive organs, injection into immature embryos, high velocity projectile introduction and the like. The choice of technique will 30 depend upon the target plant to be transformed. For example, dicotyledonous plants and certain monocots and gymnosperms may be transformed by *Agrobacterium* Ti plasmid

technology, as described, for example by Bevan (*Nucleic Acids Res.* 12:8711-8721, 1984). Targets for the introduction of the genetic constructs of the present invention include tissues, such as leaf tissue, disseminated cells, protoplasts, seeds, embryos, meristematic regions; cotyledons, hypocotyls, and the like. The preferred method for transforming eucalyptus and pine is a biolistic method using pollen (*see, for example, Aronen, Finnish Forest Res. Papers*, 595:53, 1996) or easily regenerable embryonic tissues.

Once the cells are transformed, cells having the inventive genetic construct incorporated in their genome may be selected by means of a marker, such as the kanamycin resistance marker discussed above. Transgenic cells may then be cultured in an appropriate medium to regenerate whole plants, using techniques well known in the art. In the case of protoplasts, the cell wall is allowed to reform under appropriate osmotic conditions. In the case of seeds or embryos, an appropriate germination or callus initiation medium is employed. For explants, an appropriate regeneration medium is used. Regeneration of plants is well established for many species. For a review of regeneration of forest trees, *see Dunstan et al.*, "Somatic embryogenesis in woody plants," in Thorpe TA, ed., *In vitro embryogenesis of plants*, (Current Plant Science and Biotechnology in Agriculture), Vol. 20, Chapter 12, pp. 471-540, 1995. Specific protocols for the regeneration of spruce are discussed by Roberts et al., ("Somatic embryogenesis of spruce," in Redenbaugh K, ed., *Synseed: applications of synthetic seed to crop improvement*, Chapter 23, pp. 427-449, CRC Press: [n.p.], 1993). The resulting transformed plants may be reproduced sexually or asexually, using methods well known in the art, to give successive generations of transgenic plants.

As discussed above, the production of RNA in target plant cells can be controlled by choice of the promoter sequence, or by selecting the number of functional copies or the site of integration of the polynucleotides incorporated into the genome of the target plant host. A target plant may be transformed with more than one genetic construct of the present invention, thereby modulating the activity of more than one cell wall polysaccharide enzyme, affecting enzyme activity in more than one tissue, or affecting enzyme activity at more than one expression time. Similarly, a genetic construct may be assembled containing more than one open reading frame coding for a polypeptide encoded by a polynucleotide of the present invention or more than one non-coding region of a gene coding for such a polypeptide. The polynucleotides of the present inventive may also be employed in combination with other

known sequences encoding polypeptides involved in the synthesis of cell wall polysaccharides. In this manner, it may be possible to modify a biosynthetic pathway of cell wall polysaccharides in a non-woody plant to produce a new type of woody plant.

The following examples are offered by way of illustration and not by way of  
5 limitation.

### Example 1

#### Isolation and Characterization of cDNA Clones from *Eucalyptus grandis*

*Eucalyptus grandis* cDNA expression libraries (from various tissues, including  
10 flowers, leaves, phloem, roots, seeds, shoot buds and xylem) were constructed and screened as follows.

mRNA was extracted from the plant tissue using the protocol of Chang et al. (*Plant Molecular Biology Reporter* 11:113-116, 1993) with minor modifications. Specifically, samples were dissolved in CPC-RNAXB (100 mM Tris-Cl, pH 8.0; 25 mM EDTA; 2.0 M  
15 NaCl; 2%CTAB; 2% PVP and 0.05% Spermidine\*3HCl) and extracted with chloroform:isoamyl alcohol, 24:1. mRNA was precipitated with ethanol and the total RNA preparate was purified using a Poly(A) Quik mRNA Isolation Kit (Stratagene, La Jolla, CA). A cDNA expression library was constructed from the purified mRNA by reverse transcriptase synthesis followed by insertion of the resulting cDNA clones in Lambda ZAP using a ZAP  
20 Express cDNA Synthesis Kit (Stratagene), according to the manufacturer's protocol. The resulting cDNAs were packaged using a Gigapack II Packaging Extract (Stratagene) employing 1 µl of sample DNA from the 5 µl ligation mix. Mass excision of the library was done using XL1-Blue MRF' cells and XL0LR cells (Stratagene) with ExAssist helper phage (Stratagene). The excised phagemids were diluted with NZY broth (Gibco BRL,  
25 Gaithersburg, MD) and plated out onto LB-kanamycin agar plates containing X-gal and isopropylthio-beta-galactoside (IPTG).

Of the colonies plated and picked for DNA miniprep, 99% contained an insert suitable for sequencing. Positive colonies were cultured in NZY broth with kanamycin and cDNA was purified by means of alkaline lysis and polyethylene glycol (PEG) precipitation. Agarose gel  
30 at 1% was used to screen sequencing templates for chromosomal contamination. Dye primer

sequences were prepared using a Turbo Catalyst 800 machine (Perkin Elmer/Applied Biosystems Division, Foster City, CA) according to the manufacturer's protocol.

DNA sequences for positive clones were obtained using a Perkin Elmer/Applied Biosystems Division Prism 377 sequencer. cDNA clones were sequenced first from the 5' end and, in some cases, also from the 3' end. For some clones, internal sequence was obtained using subcloned fragments. Subcloning was performed using standard procedures of restriction mapping and subcloning to pBluescript II SK+ vector.

The determined cDNA sequences are provided in SEQ ID NO: 2, 3, 6, 7, 9, 12-15, 18, 19, 21, 23, 26, 28, 29, 57, 58, 60-66, 71-73, 78, 79, 105, 107, 119-128, 139, 141, 142, 149-161, 186-195, 197, 198, 205-233, 252-256, 264, 273-293, 319-330, 366, 373-377, 382-385, 390-393, 399-434, 479-503, 507-512, 522-528, 531-547, 552-554, 556-573, 587-589, 592-612 and 621-771.

### Example 2

#### Isolation and Characterization of cDNA Clones from *Pinus radiata*

##### Isolation of cDNA clones by high through-put screening

*Pinus radiata* cDNA expression libraries (from various tissues, including cell cultures, fascicle meristems, phloem, pollen sacs, roots, seedlings, shoot buds, strobilus and xylem) were constructed and screened as described above in Example 1. DNA sequence for positive clones was obtained using forward and reverse primers on a Perkin Elmer/Applied Biosystems Division Prism 377 sequencer. The determined cDNA sequences are provided in SEQ ID NO: 1, 4, 5, 8, 10, 11, 16, 17, 20, 22, 24, 25, 27, 59, 67-70, 74-77, 80, 109-113, 140, 143, 162-185, 196, 199-204, 234-251, 257-263, 265-272, 294-318, 331-365, 367-372, 378-381, 386-389, 394-398, 435-481, 504-506, 513-521, 529, 530, 548-551, 555, 574-586, 590, 591, 613-620 and 772-908.

### Example 3

#### Polynucleotide and Amino Acid Analysis

The determined cDNA sequences described above were compared to and aligned with known sequences in the EMBL database (as updated to May 1999). Specifically, the

polynucleotides identified in SEQ ID NO: 1-29, 57-80, 105, 107, 109-111, 115-125, 135-139 and 145-904 were compared to polynucleotides in the EMBL database using the BLASTN algorithm Version 2.0.6 [Sep-16-1998] set to the following running parameters: Unix running command: blastall -p blastn -d embldb -e 10 -G0 -E0 -r1 -v30 -b30 -i queryseq -o results.

5 Multiple alignments of redundant sequences were used to build up reliable consensus sequences. Based on similarity to known sequences from other plant or non-plant species, the isolated polynucleotides of the present invention identified as SEQ ID NO: 1-29, 57-80, 105, 107, 109-113, 119-129, 139-143 and 149-908 were putatively identified as encoding the enzymes shown in Table 1, above.

10 The cDNA sequences of SEQ ID NO: 58, 60, 62, 64, 65, 67-70, 72, 74, 75, 77, 78, 80, 105, 107, 119-121, 123-128 and 139-143 were determined to have less than 40% identity to sequences in the EMBL database using the computer algorithm BLASTN, as described above. The cDNA sequences of SEQ ID NO: 57, 59, 66, 79 and 122 were determined to have less than 60% identity to sequences in the EMBL database using BLASTN, as described above.

15 The cDNA sequences of SEQ ID NO: 61, 71, 73 and 76 were determined to have less than 75% identity to sequences in the EMBL database using BLASTN, as described above. The cDNA sequence of SEQ ID NO: 63 was determined to have less than 90% identity to sequences in the EMBL database using BLASTN, as described above.

20

#### Example 4

##### Functional Identification of Cellulose Biosynthetic Genes

Sense constructs containing sequences including the coding regions for UGP (SEQ ID NO: 23) and SUS (SEQ ID NO: 49) from *Eucalyptus grandis*, and UGP (SEQ ID NO: 24) from *Pinus radiata* were inserted into the expression vector pET16b (Clontech Laboratories Inc, Palo Alto, CA). The resulting constructs were transformed into *E. coli* XL1-Blue (Stratagene) and induced to produce recombinant protein by the addition of IPTG. Purified proteins were obtained using Ni<sup>2+</sup> column chromatography (Janknecht et al., *Proc. Natl. Acad. Sci. USA*, 88:8972-8976, 1991). Enzyme assays for each of the purified proteins demonstrated the expected substrate specificity and enzymatic activity for the genes tested.

30 Enzyme assays for UGP were performed using published methods (Peng and Chang, *FEBS Lett.* 329[1,2]:153-158, 1993). The data shown in Table 2 demonstrates enzyme



activity for the expressed proteins as compared to data from Katsube et al. (*Biochem.* 30:8546-8551, 1991) and Nakano et al. (*J. Biochem.* 106:528-532, 1989).

TABLE 2

5

SEQ ID NO:	24	23	Katsube et al.	Nakano et al.
Species	<i>P. radiata</i>	<i>E. grandis</i>	<i>S. tuberosum</i>	<i>S. tuberosum</i>
Enzyme	UGP	UGP	UGP	UGP
$K_M^{GIP}$	0.121	0.126	0.130	0.180
SEM	0.020	0.002	n.a.	n.a.
$K_M^{UTP}$	0.091	not done	0.076	0.170
SEM	0.015	not done	n.a.	n.a.
$K_M^{ATP}$	no activity	no activity	no activity	no activity

Enzyme assays for SUS (sucrose synthase) were performed using the methods described by Šebková, V. et al. (*Plant Physiol.*, 108:75-83, 1995). The data shown in Table 3 demonstrates enzyme activity for the expressed proteins. The  $K_M^{Sucrose}$  of *E. grandis* is compared with the data reported by Delmer DP (*J. Biol. Chem.* 247:3822-3828, 1972) and Nakai et al. (*BioSci. Biotech. Biochem.* 61:1500-1503).

10

TABLE 3

SEQ ID NO:	49	Delmer et al.	Nakai et al.
Species	<i>E. grandis</i>	<i>V. radiata</i>	<i>V. radiata</i>
Enzyme	SUS	SUS	SUS
$K_M^{Sucrose}$	1.651	16.700	161.000
SEM	0.371	n.a.	n.a.
$K_M^{UDP}$	0.028	n.a.	n.a.
SEM	0.003	n.a.	n.a.

A sense construct containing the sequence of the coding region for cellulose synthase (CEL; SEQ ID NO: 50) from *Eucalyptus grandis* was inserted into the protein expression vector pcDNA3 (Invitrogen, Carlsbad, CA). The resulting construct was transfected into mammalian 293T cells (DuBridge RB et al., *Mol. Cell. Biol.* 7[1]:379-387, 1987), and recombinant protein was induced by the addition of IPTG. Proteins were solubilised from membranes, and the level of CEL activity was determined as described by Kudlicka K and Brown RM Jr. (*Plant Phys.* 115:643-656, 1997). As a positive control for activity, native CEL enzyme was solubilised from mung bean (*Vigna radiata*) plants. The determined levels of CEL activity for *V. radiata* are shown in Fig. 1. The levels of CEL activity found in mammalian 293T cells transfected with the *Eucalyptus* CEL expression clone were found to be similar to those obtained from *V. radiata* (Fig. 2). CEL activity was absent in non-transfected control 293T cells.

#### Example 5

##### Use of a Cellulose Synthase (CEL) Gene to Modify Polysaccharide Biosynthesis

Transformation of tobacco plants with a *Pinus radiata* CEL gene is performed as follows. Genetic constructs comprising sense and anti-sense constructs containing a polynucleotide including the coding region of CEL (SEQ ID NO: 8) from *Pinus radiata* are constructed and inserted into *Agrobacterium tumefaciens* by direct transformation using published methods (See, An G, Ebert PR, Mitra A, Ha SB, "Binary Vectors," in Gelvin SB and Schilperoort RA, eds., *Plant Molecular Biology Manual*, Kluwer Academic Publishers: Dordrecht, 1988). The constructs of sense polynucleotides are made by cloning PBK-CMV plasmid cDNA inserts into pART7 plasmids, followed by cloning of the *NotI*-digested 35S-Insert-OCS 3'UTR-fragments from the pART7 vectors into pART27 plant expression vectors (See Gleave A, "A versatile binary vector system with a T-DNA organizational structure conducive to efficient integration of cloned DNA into the plant genome," *Plant Molecular Biology* 20:1203-1207, 1992). The presence and integrity of the transgenic constructs are verified by restriction digestion and DNA sequencing.

Tobacco (*Nicotiana tabacum* cv. Samsun) leaf sections are transformed with the sense and anti-sense CEL constructs using the method of Horsch et al. (*Science* 227:1229-1231,

1985). Transformed plants containing the appropriate CEL construct are verified using Southern blot experiments. Expression of *Pinus* CEL in transformed plants is confirmed by isolating total RNA from each independent transformed plant line created with the CEL sense and anti-sense constructs. The RNA samples are analysed in Northern blot experiments to  
5 determine the level of expression of the transgene in each transformed line.

The total activity of CEL enzyme, encoded by the *Pinus* CEL gene and by the endogenous tobacco CEL gene, is analysed for each transformed plant line created with the CEL sense and anti-sense constructs. Crude protein extracts are prepared from each transformed plant and assayed using the methods of Robertson et al. (*Biochem J.*  
10 306:745-750, 1995) and Pear et al. (*Proc. Natl. Acad. Sci. USA* 93:12637-12642, 1996).

The concentration of cellulose in the transformed tobacco plants is determined using the method of Smith and Harris (*Plant Phys.* 107:1399-1409, 1995). Briefly, whole tobacco plants, of an average age of 38 days, are frozen in liquid nitrogen and ground to a fine powder in a mortar and pestle. The cellulose content of 100 mg of frozen powder from an empty  
15 vector-transformed control plant line, at least one independent transformed plant line containing the sense construct for CEL and at least one independent transformed plant lines containing the anti-sense construct for CEL are determined using a glucan estimation kit from Megazyme (Warriewood, New South Wales, Australia) using the protocols supplied by the manufacturer.

20 SEQ ID NOS: 1-908 are set out in the attached Sequence Listing. The codes for nucleotide and amino acid sequences used in the attached Sequence Listing, including the symbols "n" and "Xaa", conform to WIPO Standard ST.25 (1998), Appendix 2, Table 1.

Although the present invention has been described in some detail by way of illustration and example for purposes of clarity of understanding, changes and modifications  
25 can be carried out without departing from the scope of the invention which is intended to be limited only by the scope of the claims.

Claims:

1. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of: (1) sequences recited in SEQ ID NOS: 1-29, 57-80, 105, 107, 109-113, 119-129, 139-143 and 149-908; (2) complements of the sequences recited in SEQ ID NOS: 1-29, 57-80, 105, 107, 109-113, 119-129, 139-143 and 149-908; (3) reverse complements of the sequences recited in SEQ ID NOS: 1-29, 57-80, 105, 107, 109-113, 119-129, 139-143 and 149-908; (4) reverse sequences of the sequences recited in SEQ ID NOS: 1-29, 57-80, 105, 107, 109-113, 119-129, 139-143 and 149-908; (5) nucleotide sequences producing an Expectation ("E") value of 0.01 or less when compared to a sequence recited in (1) – (4) above; (6) nucleotide sequences having at least 50% identity to a nucleotide sequence recited in (1) – (4) above; (7) nucleotide sequences that hybridize to a sequence recited in (1) – (4) above under stringent hybridization conditions; (8) nucleotide sequences that are 200-mers of a sequence recited in (1) – (4) above; (9) nucleotide sequences that are 100-mers of a sequence recited in (1) – (4) above; (10) nucleotide sequences that are 40-mers of a sequence recited in (1) – (4) above; (11) nucleotide sequences that are 20-mers of a sequence recited in (1) – (4) above; (12) nucleotide sequences that are degeneratively equivalent to a sequence recited in (1) – (4) above; and (13) nucleotide sequences that are allelic variants of a sequence recited in (1) – (4) above.
2. An isolated oligonucleotide probe or primer comprising at least 10 contiguous residues complementary to 10 contiguous residues of a nucleotide sequence recited in claim 1.
3. A kit comprising a plurality of oligonucleotide probes or primers of claim 2.
4. A storage medium having recorded thereon a plurality of polynucleotides, at least one of the polynucleotides comprising a nucleotide sequence recited in claims 1 or 2.
5. A construct comprising a polynucleotide of claim 1.
6. A transgenic cell comprising a construct according to claim 5.
7. A construct comprising, in the 5'-3' direction:
  - (a) a gene promoter sequence;
  - (b) a polynucleotide sequence comprising at least one of the following: (1) a polynucleotide coding for at least a functional portion of a polypeptide

encoded by a nucleotide sequence of claim 1; and (2) a polynucleotide comprising a non-coding region of a gene coding for a polypeptide encoded by a nucleotide sequence selected from the group consisting of sequences recited in claim 1; and

- 5 (c) a gene termination sequence.
8. The construct of claim 7, wherein the polynucleotide is in a sense orientation.
9. The construct of claim 7, wherein the polynucleotide is in an antisense orientation.
10. The construct of claim 7, wherein the gene promoter sequence is functional in a plant host to provide for transcription in xylem.
- 10 11. A transgenic plant cell comprising a construct of claim 7.
12. A plant comprising a transgenic plant cell according to claim 11, or a part or propagule or progeny thereof.
13. A method for modulating one or more of the polysaccharide content, the polysaccharide composition and the polysaccharide structure of a plant, comprising  
15 stably incorporating into the genome of the plant a polynucleotide of claim 1.
14. The method of claim 13 wherein the plant is selected from the group consisting of eucalyptus and pine species.
15. The method of claim 13 comprising stably incorporating into the genome of the plant a construct of claim 7.
- 20 16. A method for producing a plant having one or more of altered polysaccharide content, altered polysaccharide composition and altered polysaccharide structure, comprising:
  - (a) transforming a plant cell with a construct of claim 7 to provide a transgenic cell; and
  - (b) cultivating the transgenic cell under conditions conducive to regeneration and  
25 mature plant growth.
17. A method for modifying the activity of a polypeptide involved in a polysaccharide biosynthetic pathway in a plant comprising stably incorporating into the genome of the plant a construct of claim 7.
18. An isolated polypeptide comprising an amino acid sequence selected from the group  
30 consisting of: (a) sequences of SEQ ID NOS: 30-56, 81-104, 106, 108, 114-118, 129-138 and 144-148; (b) sequences having at least 50% identity to a sequence of (a);

sequences having at least 70% identity to a sequence of (a); and sequences having at least 90% identity to a sequence of (a).

19. An isolated polypeptide encoded by an isolated polynucleotide sequence of claim 1.

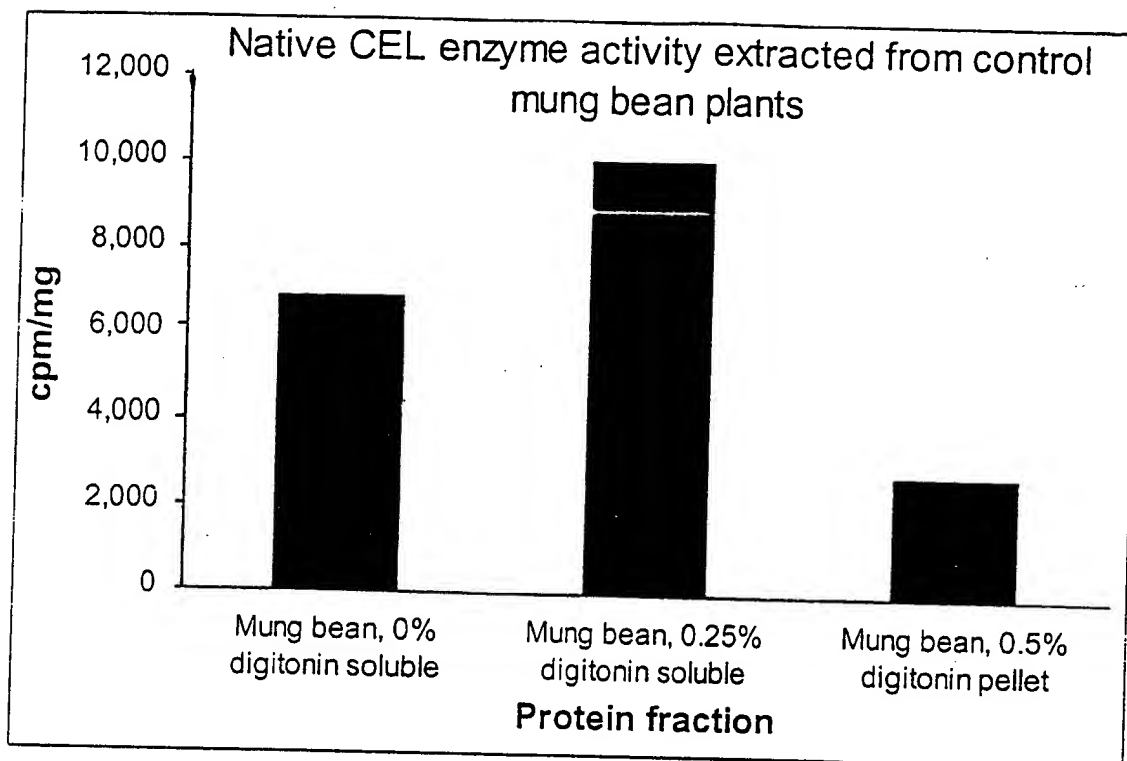
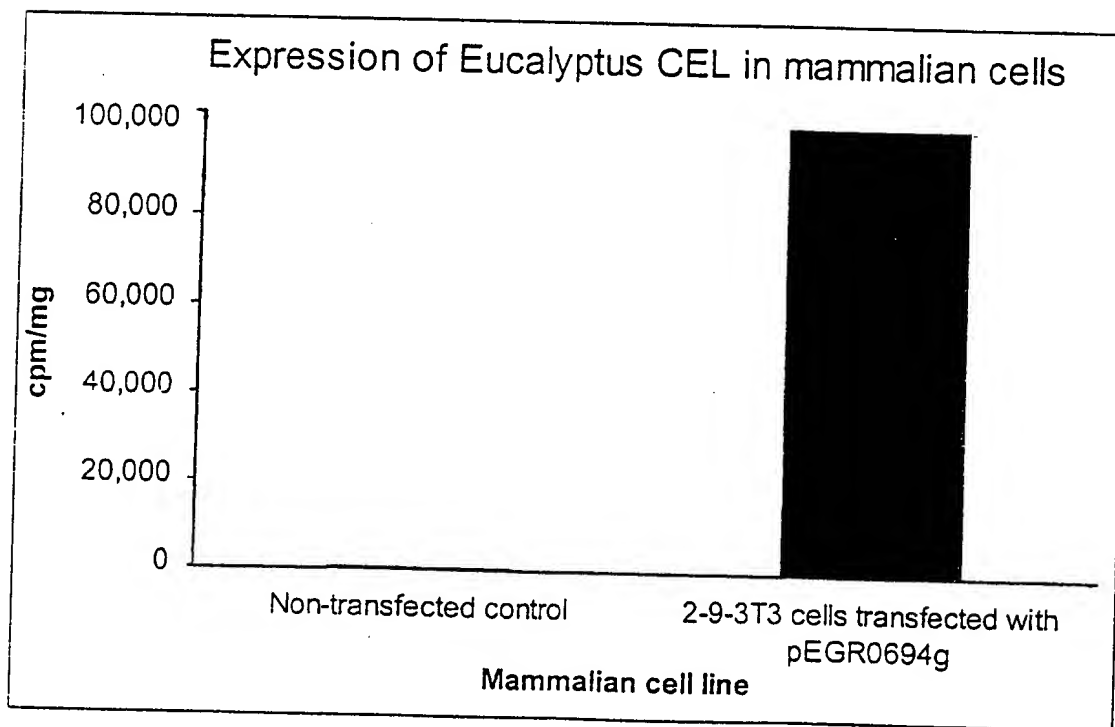
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Figure 1

Figure 2



## SEQUENCE LISTING

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<120> Materials and Methods for the  
Modification of Plant Cell Wall Polysaccharides

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<150> US 09/170,862

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&lt;210&gt; 14

&lt;211&gt; 1407

&lt;212&gt; DNA

<213> *Eucalyptus grandis*

&lt;400&gt; 14

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&lt;210&gt; 15

&lt;211&gt; 2913

&lt;212&gt; DNA

<213> *Eucalyptus grandis*

&lt;400&gt; 15

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<210> 16  
 <211> 401  
 <212> DNA  
 <213> Pinus radiata

<400> 16						
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ctaatactcc	gggtgccctt	cagaacagaa	aaaggagttt	tgcgtaactg	ggtttctcga	180
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tctttgattg	cgcataaaca	aggaatcaca	cagtgcaca	tagcccatgc	cttgaggaga	360
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<210> 17  
 <211> 477  
 <212> DNA  
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<210> 18  
 <211> 503  
 <212> DNA  
 <213> Eucalyptus grandis

<400> 18						
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<210> 19  
 <211> 413  
 <212> DNA  
 <213> Eucalyptus grandis

&lt;400&gt; 19

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tgaatggtaa	ctttgtgctt	gagcttgact	ttgagccatt	caactgcctct	tttccgcgcc	360
cgactctttc	caagtctatt	ggcaatggcg	tcgagtttctg	caatcgccat	ctc	413

&lt;210&gt; 20

&lt;211&gt; 1108

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 20

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catccgaaat	tttcgtgact	gcacatcttc	ttcgcataat	tcgccggaat	atactctcat	120
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aaagttaaaa	cttaaaattg	tgtatagc				1108

&lt;210&gt; 21

&lt;211&gt; 559

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 21

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&lt;210&gt; 22

&lt;211&gt; 1036

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

## &lt;400&gt; 22

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## &lt;210&gt; 23

&lt;211&gt; 467

&lt;212&gt; DNA

<213> *Eucalyptus grandis*

## &lt;400&gt; 23

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aagttaaccc	acaagttggt	tgtattgaaa	atcttgaact	tctcaat		467

## &lt;210&gt; 24

&lt;211&gt; 704

&lt;212&gt; DNA

<213> *Pinus radiata*

## &lt;400&gt; 24

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## &lt;210&gt; 25

&lt;211&gt; 712

&lt;212&gt; DNA



<213> *Pinus radiata*

&lt;400&gt; 25

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&lt;210&gt; 26

&lt;211&gt; 789

&lt;212&gt; DNA

<213> *Eucalyptus grandis*

&lt;400&gt; 26

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gagaagattc	acgagaaggc	ttacaatcat	gatgagctca	tcagaattgt	tactacaaga	660
agtaaagctc	agcttaatgc	aaccctcaat	tactacaaca	atgagtttgg	gaatgccatc	720
aacaaggatc	tgaaggctga	tccaaatgat	gaatttctga	aactgctgag	atcagcaatt	780
aagtgccttg						789

&lt;210&gt; 27

&lt;211&gt; 2132

&lt;212&gt; DNA

<213> *Pinus radiata*

&lt;400&gt; 27

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tgtgtttgca	cgggtcatct	gagatatgtt	gggagaagag	atcaaatcag	gtcttgtgtg	300
aagagatgtc	aacaattata	gtgccagcgc	cagctccatc	cccagttgaa	gactctgaac	360
gcctgaggaa	ggcctttgaa	ggctggggca	caaatgagaa	gttgattata	gaaatttttg	420
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gtggagtcca	aataacaggt	cacttctgga	aatctctagt	gctcgatctt	caactgagct	660
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&lt;210&gt; 28

&lt;211&gt; 2588

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 28

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gaacgcaccc	acacgatctt	ccattccctc	aacaatgtcg	actctcaccg	tcccgcagcc	120
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caataaggat	tttggctact	aggagcaaa	cacaggtcaa	tgctacgctg	aatcactaca	720
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ttgggtcata	ggaatgcggc	ctgattccgc	aaacctatgc	cgagacttac		1560
ggcgaggacc	tcctcaaggc	attggacaga	gaacttacca	atgatttcga	gaggctggtg	1620
gtcctttggt	cacttgatcc	ggctgaacgt	gatgcgtact	tggcgaatga	agcgacgaaa	1680

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aagatctcag agaaggctta tggccatgag gatctcataa ggattttggc tactaggagc 1980
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tgtcagttgt agctatgcga agcaaataca cttcttataa tggcggttgg ttatgtactt 2520
atgagaagtc tttgattttg atctttaatc aagactgcta gtaagtgatc gtgaaaaaaaa 2580
aaaaaaaaa 2588

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<210> 29  
 <211> 627  
 <212> DNA  
 <213> *Eucalyptus grandis*

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<400> 29
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agctccggaa agccttcgca ggatggggaa caaatgagaa gttgatcata tccatattgg 120
gccataggaa tgcggcgag aggaagctga ttcggcaaac ctatgccgag acttacggcg 180
aggacctcct caaggcattg gacagagaac ttaccaatga tttcgaggtc tgatcttcct 240
ttaatttttt ggattcatcc catggaagac gtgtccttct ttctctcaga ttaatccata 300
ttcattccgt atcgtcagag gctggtggtc ctttgggtcac ttgatccggc tgaacgtgat 360
gcgactttgg cgaatgaagc gacgaaaaga tggacttcaa gcaaccagg tctcatggaa 420
atagcctgca cgaggtctcc acagcagttg ctcatggcaa gacaagcata tcatgctcga 480
tacaagaagt cgctggaaga ggacgtcgct caccacacaa ctggagattt tcgtaagttg 540
taggtacctc ttgtgagctc ctaccattat gatggagatg aggtgaatat gactttggga 600
aaagcagagg ctaagatact ccacgag 627

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<210> 30  
 <211> 151  
 <212> PRT  
 <213> *Pinus radiata*

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<400> 30
Phe Arg Gly Ile Gln Tyr Arg Arg Asn Gly Trp Arg Tyr Gly Arg Arg
1      5      10      15
Val Ala Asn Leu Asn Val Leu Gly Arg Glu Thr Ala Glu Phe Thr Ser
20     25     30
Phe Arg Pro Val Phe Leu Arg Gly Asn Ser Gln Gly Leu Ser Ser Ala
35     40     45
Ser Ser Leu Cys Asp Tyr Arg Ile Phe Ala Asp Ser Lys Arg Lys Lys
50     55     60
His Ala Ile Phe Arg Lys Gln Asn Ile Asn Arg Ser Thr Val Val Ser
65     70     75     80
Pro Arg Ala Val Ser Asp Thr Phe Ser Glu Leu Thr Cys Leu Asp Pro
85     90     95
Val Ala Ser Arg Ser Val Leu Gly Ile Ile Leu Gly Gly Gly Ala Gly
100    105    110
Thr Arg Leu Tyr Pro Leu Thr Lys Lys Arg Ala Lys Pro Ala Val Pro
115    120    125

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Leu Gly Ala Asn Tyr Arg Leu Ile Asp Ile Pro Val Ser Asn Cys Ile  
 130 135 140  
 Asn Ser Asn Ile Ser Lys Ile  
 145 150

<210> 31  
 <211> 72  
 <212> PRT  
 <213> Eucalyptus grandis

<400> 31  
 Ala Pro Ala Leu Ala Ser Gly Ala Ala Ala Phe Lys Ser Val Arg Arg  
 1 5 10 15  
 Ala Pro Ala Val Val Ser Pro Arg Ala Val Ser Asp Ser Arg Asn Ser  
 20 25 30  
 Gln Thr Cys Leu Asp Pro Asp Ala Ser Arg Ser Val Leu Gly Ile Ile  
 35 40 45  
 Leu Gly Gly Gly Ala Gly Thr Arg Leu Tyr Pro Leu Thr Lys Lys Arg  
 50 55 60  
 Ala Lys Pro Ala Val Pro Leu Gly  
 65 70

<210> 32  
 <211> 124  
 <212> PRT  
 <213> Eucalyptus grandis

<400> 32  
 Leu Lys Asp Ala Ile Ile Ser His Gly Cys Phe Leu Arg Glu Cys Arg  
 1 5 10 15  
 Val Glu Arg Ser Ile Val Gly Glu Arg Ser Arg Leu Asp Ser Gly Val  
 20 25 30  
 Glu Leu Lys Asp Thr Val Met Met Gly Ala Asp Tyr Tyr Gln Thr Glu  
 35 40 45  
 Ser Glu Ile Ala Ser Leu Leu Ala Glu Gly Lys Val Pro Ile Gly Ile  
 50 55 60  
 Gly Lys Asn Thr Lys Ile Arg Asn Cys Ile Ile Asp Lys Asn Ala Lys  
 65 70 75 80  
 Ile Gly Lys Asp Val Ala Ile Val Asn Lys Asp Gly Val Glu Glu Ala  
 85 90 95  
 Asp Arg Pro Gly Asp Gly Phe Tyr Ile Arg Leu Gly Ile Thr Val Ile  
 100 105 110  
 Leu Glu Lys Ala Thr Ile Glu Asp Gly Thr Val Ile  
 115 120

<210> 33  
 <211> 67  
 <212> PRT  
 <213> Pinus radiata

<400> 33  
 Ile Arg Ile Val Leu Gln Gly Phe Asn Trp Glu Ser His Arg Ser Gly  
 1 5 10 15  
 Gly Trp Tyr His Lys Leu Ser Gly Lys Ala Ala Glu Ile Ala Ser Lys  
 20 25 30  
 Gly Phe Thr Ile Val Trp Leu Pro Pro Pro Thr Asp Ser Val Ser Pro  
 35 40 45

Glu Gly Tyr Met Pro Arg Asp Leu Tyr Asp Leu Asn Ser Arg Tyr Gly  
 50 55 60  
 Ser Leu Glu  
 65

<210> 34  
 <211> 157  
 <212> PRT  
 <213> Pinus radiata

<400> 34  
 Asn Gln Asp Ala His Arg Gln Arg Ile Val Asn Trp Ile Asn Ala Thr  
 1 5 10 15  
 Gly Gly Ser Ser Ser Ala Phe Asp Val Thr Thr Lys Gly Ile Leu His  
 20 25 30  
 Val Ala Leu His Asn Gln Tyr Trp Arg Leu Ile Asp Pro Gln Gly Lys  
 35 40 45  
 Pro Thr Gly Val Met Gly Trp Trp Pro Ser Arg Ala Val Thr Tyr Leu  
 50 55 60  
 Glu Asn His Asp Thr Gly Ser Thr Gln Gly His Trp Pro Phe Pro Arg  
 65 70 75 80  
 Asp Lys Leu Thr Gln Gly Tyr Ala Tyr Ile Leu Thr His Pro Gly Thr  
 85 90 95  
 Pro Thr Ile Phe Tyr Asp His Phe Tyr Asp Phe Gly Leu His Asp Thr  
 100 105 110  
 Ile Thr Glu Leu Ile Asp Ala Arg Thr Arg Ala Gly Ile His Cys Arg  
 115 120 125  
 Ser Thr Leu Lys Ile Phe His Ala Asn Asn Glu Gly Tyr Ala Ala Gln  
 130 135 140  
 Ile Asp Glu Asn Leu Val Met Lys Leu Gly Gln Phe Asp  
 145 150 155

<210> 35  
 <211> 332  
 <212> PRT  
 <213> Eucalyptus grandis

<400> 35  
 Pro Thr Asp Ser Val Ser Pro Glu Gly Tyr Met Pro Arg Asp Leu Tyr  
 1 5 10 15  
 Asn Leu Asn Ser Arg Tyr Gly Thr Ile Asp Glu Leu Lys Asp Leu Val  
 20 25 30  
 Lys Lys Phe His Glu Val Asn Ile Arg Val Leu Gly Asp Val Val Leu  
 35 40 45  
 Asn His Arg Cys Ala Gln Tyr Gln Asn Gln Asn Gly Ile Trp Asn Ile  
 50 55 60  
 Phe Gly Gly Arg Leu Asn Trp Asp Asp Arg Ala Val Val Ala Asp Asp  
 65 70 75 80  
 Pro His Phe Gln Gly Arg Gly Asn Lys Ser Ser Gly Asp Asn Phe His  
 85 90 95  
 Ala Ala Pro Asn Ile Asp His Ser Gln Asp Phe Val Arg Lys Asp Leu  
 100 105 110  
 Lys Glu Trp Leu His Trp Leu Arg Ser Glu Ile Gly Tyr Asp Gly Trp  
 115 120 125  
 Arg Leu Asp Phe Val Arg Gly Phe Trp Gly Gly Tyr Val Lys Asp Tyr  
 130 135 140  
 Leu Asp Ala Ser Glu Pro Tyr Phe Ala Val Gly Glu Tyr Trp Asp Ser

145                      150                      155                      160  
 Leu Ser Tyr Thr Tyr Gly Glu Met Asp His Asn Gln Asp Ala His Arg  
                                  165                      170                      175  
 Gln Arg Ile Ile Asp Trp Ile Asn Ala Thr Asn Gly Thr Ala Gly Ala  
                                  180                      185                      190  
 Phe Asp Val Thr Thr Lys Gly Ile Leu His Ala Ala Leu Glu Arg Cys  
                                  195                      200                      205  
 Glu Tyr Trp Arg Leu Ser Asp Gln Lys Gly Lys Pro Pro Gly Val Val  
                                  210                      215                      220  
 Gly Trp Trp Pro Ser Arg Ala Val Thr Phe Val Glu Asn His Asp Thr  
 225                                   230                      235                      240  
 Gly Ser Thr Gln Gly His Trp Arg Phe Pro Ser Gly Lys Glu Met Gln  
                                  245                      250                      255  
 Gly Tyr Ala Tyr Ile Leu Thr His Pro Gly Thr Pro Ala Val Phe Tyr  
                                  260                      265                      270  
 Asp His Ile Phe Ser His Tyr Gln Ser Glu Ile Gly Ser Leu Ile Ser  
                                  275                      280                      285  
 Ile Arg Asn Arg Asn Lys Ile His Cys Arg Ser Thr Ile Lys Ile Thr  
                                  290                      295                      300  
 Lys Ala Glu Arg Asp Val Tyr Ala Ala Ile Ile Asp Asp Lys Val Ala  
 305                                   310                      315                      320  
 Met Lys Ile Gly Pro Gly Tyr Tyr Glu Pro Gln Ser  
                                  325                      330

&lt;210&gt; 36

&lt;211&gt; 251

&lt;212&gt; PRT

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 36

Met Met Glu Ser Gly Val Pro Leu Cys Asn Thr Cys Gly Glu Ala Val  
 1                                   5                                   10                                   15  
 Gly Val Asp Glu Lys Gly Glu Val Phe Val Ala Cys Gln Glu Cys Asn  
                                  20                                   25                                   30  
 Phe Ala Ile Cys Lys Ala Cys Val Glu Tyr Glu Ile Lys Glu Gly Arg  
                                  35                                   40                                   45  
 Lys Ala Cys Leu Arg Cys Gly Thr Pro Phe Glu Ala Asn Ser Met Ala  
                                  50                                   55                                   60  
 Asp Ala Glu Arg Asn Glu Leu Gly Ser Arg Ser Thr Met Ala Ala Gln  
 65                                   70                                   75                                   80  
 Leu Asn Asp Pro Gln Asp Thr Gly Ile His Ala Arg His Ile Ser Ser  
                                  85                                   90                                   95  
 Val Ser Thr Leu Asp Ser Glu Tyr Asn Asp Glu Thr Gly Asn Pro Ile  
                                  100                                   105                                   110  
 Trp Lys Asn Arg Val Glu Ser Trp Lys Asp Lys Lys Asn Lys Lys Lys  
                                  115                                   120                                   125  
 Lys Ala Pro Thr Lys Ala Glu Lys Glu Ala Gln Val Pro Pro Glu Gln  
                                  130                                   135                                   140  
 Gln Met Glu Glu Lys Gln Ile Ala Asp Ala Ser Glu Pro Leu Ser Thr  
 145                                   150                                   155                                   160  
 Val Ile Pro Ile Ala Lys Ser Lys Leu Ala Pro Tyr Arg Thr Val Ile  
                                  165                                   170                                   175  
 Ile Met Arg Leu Ile Ile Leu Ala Leu Phe Phe His Tyr Arg Val Thr  
                                  180                                   185                                   190  
 His Pro Val Asp Ser Ala Tyr Pro Leu Trp Leu Thr Ser Ile Ile Cys  
                                  195                                   200                                   205  
 Glu Ile Trp Phe Ala Tyr Ser Trp Val Leu Asp Gln Phe Pro Lys Trp

210                      215                      220  
 Ser Pro Val Asn Arg Ile Thr His Val Asp Arg Leu Ser Ala Arg Tyr  
 225                      230                      235                      240  
 Glu Lys Glu Gly Glu Pro Ser Glu Leu Ala Val  
                     245                      250

<210> 37  
 <211> 127  
 <212> PRT  
 <213> Pinus radiata

<400> 37  
 Leu Pro Arg Leu Val Tyr Val Ser Arg Glu Lys Arg Pro Gly Tyr Gln  
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 His His Lys Lys Ala Gly Ala Met Asn Ala Leu Val Arg Val Ser Ala  
                     20                      25                      30  
 Val Leu Thr Asn Ala Pro Phe Ile Leu Asn Leu Asp Cys Asp His Tyr  
                     35                      40                      45  
 Leu Asn Asn Ser Lys Ala Val Arg Glu Ala Met Cys Phe Leu Met Asp  
 50                      55                      60  
 Pro Gln Leu Gly Lys Lys Leu Cys Tyr Val Gln Phe Pro Gln Arg Phe  
 65                      70                      75                      80  
 Asp Gly Ile Asp Arg His Asp Arg Tyr Ala Asn Arg Asn Thr Val Phe  
                     85                      90                      95  
 Phe Asp Ile Asn Met Lys Gly Leu Asp Gly Ile Gln Gly Pro Val Tyr  
                     100                      105                      110  
 Val Gly Thr Gly Cys Val Phe Asn Arg Gln Ala Leu Tyr Gly Tyr  
                     115                      120                      125

<210> 38  
 <211> 534  
 <212> PRT  
 <213> Eucalyptus grandis

<400> 38  
 His Tyr Ile Asn Asn Ser Lys Ala Ile Arg Glu Ala Met Cys Phe Leu  
 1                      5                      10                      15  
 Met Asp Pro Gln Leu Gly Lys Lys Leu Cys Tyr Val Gln Phe Pro Gln  
                     20                      25                      30  
 Arg Phe Asp Gly Ile Asp Arg His Asp Arg Tyr Ala Asn Arg Asn Ile  
                     35                      40                      45  
 Val Phe Phe Asp Ile Asn Met Arg Gly Leu Asp Gly Ile Gln Gly Pro  
 50                      55                      60  
 Val Tyr Val Gly Thr Gly Cys Val Phe Asn Arg Gln Ala Leu Tyr Gly  
 65                      70                      75                      80  
 Tyr Asp Pro Pro Val Ser Gln Lys Arg Pro Lys Met Thr Cys Asp Cys  
                     85                      90                      95  
 Trp Pro Ser Trp Cys Ser Cys Cys Cys Gly Gly Ser Arg Lys Ser Lys  
                     100                      105                      110  
 Ser Lys Lys Lys Asp Asp Thr Ser Leu Leu Gly Pro Val His Ala Lys  
                     115                      120                      125  
 Lys Lys Lys Met Thr Gly Lys Asn Tyr Leu Lys Lys Lys Gly Ser Gly  
 130                      135                      140  
 Pro Val Phe Asp Leu Glu Asp Ile Glu Glu Gly Leu Glu Gly Phe Asp  
 145                      150                      155                      160  
 Glu Leu Glu Lys Ser Ser Leu Met Ser Gln Lys Asn Phe Glu Lys Arg  
                     165                      170                      175

Phe Gly Gln Ser Pro Val Phe Ile Ala Ser Thr Leu Met Glu Asp Gly  
 180 185 190  
 Gly Leu Pro Glu Gly Thr Asn Ser Thr Ser Leu Ile Lys Glu Ala Ile  
 195 200 205  
 His Val Ile Ser Cys Gly Tyr Glu Glu Lys Thr Glu Trp Gly Lys Glu  
 210 215 220  
 Ile Gly Trp Ile Tyr Gly Ser Val Thr Glu Asp Ile Leu Thr Gly Phe  
 225 230 235 240  
 Lys Met His Cys Arg Gly Trp Lys Ser Val Tyr Cys Met Pro Lys Arg  
 245 250 255  
 Pro Ala Phe Lys Gly Ser Ala Pro Ile Asn Leu Ser Asp Arg Leu His  
 260 265 270  
 Gln Val Leu Arg Trp Ala Leu Gly Ser Val Glu Ile Phe Leu Ser Arg  
 275 280 285  
 His Cys Pro Leu Trp Tyr Ala Trp Gly Gly Lys Leu Lys Leu Leu Glu  
 290 295 300  
 Arg Leu Ala Tyr Ile Asn Thr Ile Val Tyr Pro Phe Thr Ser Ile Pro  
 305 310 315 320  
 Leu Leu Phe Tyr Cys Thr Ile Pro Ala Val Cys Leu Leu Thr Gly Lys  
 325 330 335  
 Phe Ile Ile Pro Thr Leu Thr Asn Phe Ala Ser Ile Trp Phe Leu Ala  
 340 345 350  
 Leu Phe Leu Ser Ile Ile Ala Thr Gly Val Leu Glu Leu Arg Trp Ser  
 355 360 365  
 Gly Val Ser Ile Glu Asp Trp Trp Arg Asn Glu Gln Phe Trp Val Ile  
 370 375 380  
 Gly Gly Val Ser Ala His Leu Phe Ala Val Phe Gln Gly Leu Leu Lys  
 385 390 395 400  
 Val Leu Ala Gly Val Asp Thr Asn Phe Thr Val Thr Ala Lys Ala Ala  
 405 410 415  
 Glu Asp Ser Glu Phe Gly Glu Leu Tyr Leu Phe Lys Trp Thr Thr Leu  
 420 425 430  
 Leu Ile Pro Thr Thr Leu Ile Ile Leu Asn Met Val Gly Val Val  
 435 440 445  
 Ala Gly Val Ser Asp Ala Ile Asn Asn Gly Tyr Gly Ser Trp Gly Pro  
 450 455 460  
 Leu Phe Gly Lys Leu Phe Phe Ala Phe Trp Val Ile Val His Leu Tyr  
 465 470 475 480  
 Pro Phe Leu Lys Gly Leu Met Gly Lys Gln Asn Arg Thr Pro Thr Ile  
 485 490 495  
 Val Val Leu Trp Ser Val Leu Leu Ala Ser Ile Phe Ser Leu Val Trp  
 500 505 510  
 Val Arg Ile Asp Pro Phe Leu Pro Lys Gln Thr Gly Pro Val Leu Lys  
 515 520 525  
 Pro Cys Gly Val Glu Cys  
 530

<210> 39  
 <211> 133  
 <212> PRT  
 <213> Pinus radiata

<400> 39  
 Leu Ala Leu Phe Leu Thr Trp Arg Val Lys Asn Pro Asn Thr Asp Ala  
 1 5 10 15  
 Tyr Trp Leu Trp Gly Met Ser Ile Val Cys Glu Leu Trp Phe Ala Phe  
 20 25 30



Ser Trp Leu Leu Asp Gln Leu Pro Lys Leu Cys Pro Ile Asn Arg Ser  
 35 40 45  
 Thr Asp Leu Ala Val Leu Lys Asp Lys Phe Glu Ser Pro Thr Gly Asp  
 50 55 60  
 Asn Pro Ala Gly Arg Ser Asp Leu Pro Gly Ile Asp Cys Phe Val Ser  
 65 70 75 80  
 Thr Ala Asp Pro Glu Lys Glu Pro Pro Leu Val Thr Ala Asn Thr Ile  
 85 90 95  
 Leu Ser Ile Leu Ser Ala Asp Tyr Pro Val Glu Lys Leu Ala Cys Tyr  
 100 105 110  
 Val Ser Asp Gly Gly Ala Leu Thr Phe Glu Ala Met Ala Glu  
 115 120 125  
 Ala Ala Ser Phe Ala  
 130

<210> 40  
 <211> 206  
 <212> PRT  
 <213> Pinus radiata

<400> 40  
 Leu Leu Val Ser Gln Arg Ser Phe Glu Lys Ser Phe Gly Gln Ser Ser  
 1 5 10 15  
 Val Phe Ile Ala Ser Thr Leu Met Asp Asn Gly Gly Val Pro Glu Ser  
 20 25 30  
 Thr Asn Pro Ala Ser Leu Ile Lys Glu Ala Ile His Val Ile Ser Cys  
 35 40 45  
 Gly Tyr Glu Glu Lys Thr Glu Trp Gly Lys Glu Val Gly Trp Ile Tyr  
 50 55 60  
 Gly Ser Val Thr Glu Asp Ile Leu Thr Gly Phe Lys Met His Cys Arg  
 65 70 75 80  
 Gly Trp Arg Ser Ile Tyr Cys Met Pro Lys Arg Pro Ala Phe Lys Gly  
 85 90 95  
 Ser Ala Pro Ile Asn Leu Ser Asp Arg Leu His Gln Val Leu Arg Trp  
 100 105 110  
 Ala Leu Gly Ser Ile Glu Ile Leu Phe Ser Arg His Cys Pro Leu Trp  
 115 120 125  
 Tyr Gly Phe Gly Ala Gly Arg Leu Lys Trp Leu Glu Arg Leu Ala Tyr  
 130 135 140  
 Thr Asn Thr Ile Val Tyr Pro Leu Thr Ser Leu Pro Leu Ile Ala Tyr  
 145 150 155 160  
 Cys Thr Leu Pro Ala Ile Cys Leu Leu Thr Gly Glu Phe Ile Ile Pro  
 165 170 175  
 Thr Leu Ser Asn Leu Ala Ser Ile Tyr Phe Met Leu Leu Phe Ile Ser  
 180 185 190  
 Ile Ile Val Thr Gly Val Leu Glu Leu Arg Trp Ser Gly Val  
 195 200 205

<210> 41  
 <211> 239  
 <212> PRT  
 <213> Eucalyptus grandis

<400> 41  
 Leu Ala Leu Arg His Asp Arg Glu Gly Glu Pro Ser Gln Leu Ala Pro  
 1 5 10 15  
 Val Asp Val Phe Val Ser Thr Val Asp Pro Leu Lys Glu Pro Pro Leu

```

      20      25      30
Ile Thr Ala Asn Thr Val Leu Ser Ile Leu Ala Val Asp Tyr Pro Val
      35      40      45
Asp Lys Val Ser Cys Tyr Val Ser Asp Asp Gly Ser Ala Met Leu Thr
      50      55      60
Phe Glu Ala Leu Ser Glu Thr Ala Glu Phe Ala Arg Lys Trp Val Pro
      65      70      75      80
Phe Cys Lys Lys His Asn Ile Glu Pro Arg Ala Pro Glu Phe Tyr Phe
      85      90      95
Ala Gln Lys Ile Asp Tyr Leu Lys Asp Lys Ile Gln Pro Ser Phe Val
      100      105      110
Lys Glu Arg Arg Ala Met Lys Arg Glu Tyr Glu Glu Phe Lys Val Arg
      115      120      125
Ile Asn Ala Leu Val Ala Lys Ala Gln Lys Met Pro Glu Glu Gly Trp
      130      135      140
Thr Met Gln Asp Gly Thr Ala Trp Pro Gly Asn Asn Pro Arg Asp His
      145      150      155      160
Pro Gly Met Ile Gln Val Phe Leu Gly His Ser Gly Gly Leu Asp Thr
      165      170      175
Asp Gly Asn Glu Leu Pro Arg Leu Val Tyr Val Ser Arg Glu Lys Arg
      180      185      190
Pro Gly Phe Gln His His Lys Lys Ala Gly Ala Met Asn Ala Leu Ile
      195      200      205
Arg Val Ser Ala Val Leu Thr Asn Gly Ala Tyr Leu Leu Asn Val Asp
      210      215      220
Cys Asp His Tyr Phe Asn Asn Ser Lys Ala Leu Lys Glu Ala Met
      225      230      235

```

<210> 42  
 <211> 253  
 <212> PRT  
 <213> Eucalyptus grandis

```

      <400> 42
Ile Ser Cys Gly Tyr Glu Asp Lys Thr Glu Trp Gly Lys Glu Ile Gly
      1      5      10      15
Trp Ile Tyr Gly Ser Val Thr Glu Asp Ile Leu Thr Gly Phe Lys Met
      20      25      30
His Ala Arg Gly Trp Ile Ser Ile Tyr Cys Met Pro Pro Arg Pro Ala
      35      40      45
Phe Lys Gly Ser Ala Pro Ile Asn Leu Ser Asp Arg Leu Asn Gln Val
      50      55      60
Leu Arg Trp Ala Leu Gly Ser Ile Glu Ile Leu Leu Ser Arg His Cys
      65      70      75      80
Pro Ile Trp Tyr Gly Tyr Asn Gly Lys Leu Arg Leu Leu Glu Arg Leu
      85      90      95
Ala Tyr Ile Asn Thr Ile Val Tyr Pro Leu Thr Ser Ile Pro Leu Ile
      100      105      110
Ala Tyr Cys Ile Leu Pro Ala Phe Cys Leu Leu Thr Asn Lys Phe Ile
      115      120      125
Ile Pro Glu Ile Ser Asn Phe Ala Ser Met Trp Phe Ile Leu Leu Phe
      130      135      140
Val Ser Ile Phe Thr Thr Gly Ile Leu Glu Leu Arg Trp Ser Gly Val
      145      150      155      160
Ser Ile Glu Asp Trp Trp Arg Asn Glu Gln Phe Trp Val Ile Gly Gly
      165      170      175
Thr Ser Ala His Leu Phe Ala Val Phe Gln Gly Leu Leu Lys Val Leu

```

180										185					190				
Ala	Gly	Ile	Asp	Thr	Asn	Phe	Thr	Val	Thr	Ser	Lys	Ala	Gly	Asp	Glu				
			195			200						205							
Asp	Gly	Asp	Phe	Ala	Glu	Leu	Tyr	Val	Phe	Lys	Trp	Thr	Ser	Leu	Leu				
			210			215						220							
Ile	Pro	Pro	Thr	Thr	Val	Leu	Ile	Val	Asn	Ile	Ile	Gly	Ile	Val	Ala				
			225			230						235			240				
Gly	Val	Ser	Tyr	Ala	Ile	Asn	Ser	Gly	Tyr	Gln	Ser	Trp							
			245			250													

```
<210> 43
<211> 469
<212> PRT
<213> Eucalyptus grandis
```

<400> 43															
Gln 1	Cys	Phe	Ser	Ile 5	Leu	Ala	Val	Asp	Tyr 10	Pro	Val	Asp	Lys	Val 15	Ser
Cys	Tyr	Leu	Ser	Asp	Asp	Gly	Ala	Ala 25	Met	Leu	Ser	Phe	Glu 30	Ser	Leu
Val	Glu	Thr 35	Ala	Asp	Phe	Ala	Arg 40	Lys	Trp	Val	Pro	Phe 45	Cys	Lys	Lys
Tyr 50	Ser	Ile	Glu	Pro	Arg	Ala 55	Pro	Glu	Phe	Tyr	Phe 60	Ser	Gln	Lys	Ile
Asp 65	Tyr	Leu	Lys	Asp	Lys 70	Ile	Gln	Pro	Ser	Phe 75	Val	Lys	Glu	Arg	Arg 80
Ala	Met	Lys	Arg	Asp	Tyr 85	Glu	Glu	Phe 90	Lys	Val	Arg	Val	Asn 95	Ala	Leu
Val	Ala	Lys	Ala 100	Gln	Lys	Ala	Pro	Glu 105	Gly	Trp	Ser	Met 110	Gln	Asp	
Gly	Thr 115	Pro	Trp	Pro	Gly	Asn	Asn 120	Ser	Arg	Asp	His 125	Pro	Gly	Met	Ile
Gln 130	Val	Phe	Leu	Gly	Ser	Ser 135	Gly	Ala	His	Asp 140	Ile	Glu	Gly	Asn	Glu
Leu 145	Pro	Arg	Leu	Val	Tyr 150	Val	Ser	Arg	Glu	Lys 155	Arg	Pro	Gly	Phe	Gln 160
His	His	Lys	Lys	Ala 165	Gly	Ala	Glu	Asn 170	Ala	Leu	Val	Arg 175	Val	Ser	Ala 175
Ile	Leu	Thr 180	Asn	Ala	Pro	Tyr	Ile 185	Leu	Asn	Leu	Asp	Cys 190	Asp	His	Tyr 190
Val	Asn	Tyr 195	Ser	Asn	Ala	Val	Arg 200	Glu	Ala	Met	Cys 205	Phe	Leu	Met	Asp 205
Pro	Gln 210	Val	Gly	Arg	Asn 215	Leu	Cys	Tyr	Val	Gln 220	Phe	Pro	Gln	Arg	Phe 220
Asp 225	Gly	Ile	Asp	Arg	Ser 230	Asp	Arg	Tyr	Ala 235	Asn	Arg	Asn	Thr	Val	Phe 240
Phe	Asp	Val	Asn	Met 245	Lys	Gly	Leu	Asp 250	Gly	Ile	Gln	Gly	Pro	Val	Tyr 255
Val	Gly	Thr 260	Gly	Cys	Val	Phe	Asn 265	Arg	Gln	Ala	Leu	Tyr 270	Gly	Tyr	Gly 270
Pro	Pro 275	Ser	Met	Pro	Asn 280	Leu	Pro	Lys	Pro	Ser	Ser 285	Ser	Cys	Ser	Trp 285
Cys 290	Gly	Cys	Cys	Ser	Cys 295	Cys	Cys	Pro	Ser	Lys 300	Lys	Pro	Thr	Lys	Asp 300
Leu 305	Ser	Glu	Val	Tyr	Arg 310	Asp	Ser	Lys	Arg 315	Glu	Asp	Leu	Asn	Ala	Ala 320
Ile	Phe	Asn	Leu	Gly	Glu	Ile	Asp	Asn	Tyr	Asp	Glu	His	Glu	Arg	Ser

```

          325          330          335
Met Leu Ile Ser Gln Met Ser Phe Glu Lys Thr Phe Gly Leu Ser Thr
          340          345          350
Val Phe Ile Glu Ser Thr Leu Leu Ala Asn Gly Gly Val Pro Glu Ser
          355          360          365
Ala His Pro Ser Met Leu Ile Lys Glu Ala Ile His Val Ile Ser Cys
          370          375          380
Gly Tyr Glu Glu Lys Thr Ala Trp Gly Lys Glu Ile Gly Trp Ile Tyr
          385          390          395          400
Gly Ser Val Thr Glu Asp Ile Leu Thr Gly Phe Lys Met His Cys Arg
          405          410          415
Gly Trp Arg Ser Val Tyr Cys Met Pro Leu Arg Pro Ala Phe Lys Gly
          420          425          430
Ser Ala Pro Ile Asn Leu Ser Asp Arg Leu His Gln Val Leu Arg Trp
          435          440          445
Ala Leu Gly Ser Val Glu Ile Phe Leu Ser Arg His Cys Pro Leu Trp
          450          455          460
Tyr Gly Phe Gly Gly
          465

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<210> 44
<211> 805
<212> PRT
<213> Eucalyptus grandis

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          <400> 44
Met Ala Asp Arg Met Leu Thr Arg Ser His Ser Leu Arg Glu Arg Leu
  1          5          10          15
Asp Glu Thr Leu Ser Ala His Arg Asn Asp Ile Val Ala Phe Leu Ser
          20          25          30
Arg Val Glu Ala Lys Gly Lys Gly Ile Leu Gln Arg His Gln Ile Phe
          35          40          45
Ala Glu Phe Glu Ala Ile Ser Glu Glu Ser Arg Ala Lys Leu Leu Asp
          50          55          60
Gly Ala Phe Gly Glu Val Leu Lys Ser Thr Gln Glu Ala Ile Val Ser
          65          70          75          80
Pro Pro Trp Val Ala Leu Ala Val Arg Pro Arg Pro Gly Val Trp Glu
          85          90          95
His Ile Arg Val Asn Val His Ala Leu Val Leu Glu Gln Leu Glu Val
          100          105          110
Ala Glu Tyr Leu His Phe Lys Glu Glu Leu Ala Asp Gly Ser Leu Asn
          115          120          125
Gly Asn Phe Val Leu Glu Leu Asp Phe Glu Pro Phe Thr Ala Ser Phe
          130          135          140
Pro Arg Pro Thr Leu Ser Lys Ser Ile Gly Asn Gly Val Glu Phe Leu
          145          150          155          160
Asn Arg His Leu Ser Ala Lys Leu Phe His Asp Lys Glu Ser Leu His
          165          170          175
Pro Leu Leu Glu Phe Leu Gln Val His Cys Tyr Lys Gly Lys Asn Met
          180          185          190
Met Val Asn Ala Arg Ile Gln Asn Val Phe Ser Leu Gln His Val Leu
          195          200          205
Arg Lys Ala Glu Glu Tyr Leu Thr Ser Leu Lys Pro Glu Thr Pro Tyr
          210          215          220
Ser Gln Phe Glu His Lys Phe Gln Glu Ile Gly Leu Glu Arg Gly Trp
          225          230          235          240
Gly Asp Thr Ala Glu Arg Val Leu Glu Met Ile Gln Leu Leu Leu Asp

```

22

Lys Ser Gly Tyr His Ile Asp Pro Tyr His Gly Asp Gln Ala Ala Glu  
 705 710 715 720  
 Leu Leu Val Asp Phe Phe Asn Lys Cys Lys Ile Asp Gln Ser His Trp  
 725 730 735  
 Asp Glu Ile Ser Lys Gly Ala Met Gln Arg Ile Glu Glu Lys Tyr Thr  
 740 745 750  
 Trp Lys Ile Tyr Ser Glu Arg Leu Leu Asn Leu Thr Ala Val Tyr Gly  
 755 760 765  
 Phe Trp Lys His Val Thr Asn Leu Asp Arg Arg Glu Ser Arg Arg Tyr  
 770 775 780  
 Leu Glu Met Phe Tyr Ala Leu Lys Tyr Arg Pro Leu Ala Gln Ser Val  
 785 790 795 800  
 Pro Pro Ala Val Glu  
 805

<210> 45  
 <211> 133  
 <212> PRT  
 <213> Pinus radiata

<400> 45  
 Ile Lys Gln Gln Gly Leu Asp Ile Thr Pro Gln Ile Ile Val Val Thr  
 1 5 10 15  
 Arg Leu Ile Pro Glu Ala His Gly Thr Thr Cys Asn Gln Arg Ile Glu  
 20 25 30  
 Lys Val Ser Gly Thr Gln His Ser Leu Ile Leu Arg Val Pro Phe Arg  
 35 40 45  
 Thr Glu Lys Gly Val Leu Arg Asn Trp Val Ser Arg Phe Asp Val Trp  
 50 55 60  
 Pro Tyr Leu Glu Arg Phe Ser Glu Asp Val Thr Asn Glu Val Thr Ala  
 65 70 75 80  
 Glu Leu Lys Gly Gln Pro Asp Leu Ile Ile Gly Asn Tyr Ser Asp Gly  
 85 90 95  
 Asn Leu Val Ala Ser Leu Ile Ala His Lys Gln Gly Ile Thr Gln Cys  
 100 105 110  
 Asn Ile Ala His Ala Leu Glu Lys Thr Lys Tyr Pro Asp Ser Asp Ile  
 115 120 125  
 Tyr Trp Lys Asn Phe  
 130

<210> 46  
 <211> 158  
 <212> PRT  
 <213> Pinus radiata

<400> 46  
 His Gly Ile Asp Val Phe Asp Pro Lys Phe Asn Ile Val Ser Pro Gly  
 1 5 10 15  
 Ala Asp Met Gln Ile Tyr Phe Pro Tyr Thr Glu Lys Gln His Arg Leu  
 20 25 30  
 Thr Thr Leu His Gly Thr Ile Glu Glu Leu Leu Phe Ser Pro Glu Gln  
 35 40 45  
 Thr Ala Glu His Met Cys Ala Leu Asn Asp Arg Lys Lys Pro Ile Ile  
 50 55 60  
 Phe Ser Met Ala Arg Leu Asp Arg Val Lys Asn Met Thr Gly Leu Val  
 65 70 75 80  
 Glu Trp Phe Ala Lys Ser Lys Arg Leu Arg Glu Leu Val Asn Leu Val

```

      85          90          95
Val Val Ala Gly Asp Ile Asp Pro Ser Lys Ser Lys Asp Arg Glu Glu
      100          105          110
Val Ala Glu Ile Glu Lys Met His Arg Leu Val Lys Glu Tyr Asn Leu
      115          120          125
Asn Gly Gln Phe Arg Trp Ile Cys Ala Gln Lys Asn Arg Val Arg Asn
      130          135          140
Gly Glu Leu Tyr Arg Tyr Ile Cys Asp Thr Arg Gly Ala Phe
145          150          155

```

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<210> 47
<211> 144
<212> PRT
<213> Eucalyptus grandis

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```

<400> 47
Met Ala Asp Arg Val Leu Asn Arg Ser His Ser Pro Arg Glu Arg Leu
 1      5      10      15
Asp Glu Ala Leu Phe Ala Asp Arg Asn Asp Cys Leu Val Phe Leu Ser
      20      25      30
Arg Leu Lys Ala Lys Gly Lys Gly Ile Leu Gln Arg His Gln Ile Leu
      35      40      45
Ala Val Phe Glu Ala Ile Pro Glu Glu Ser Arg Ala Arg Leu Leu Asp
      50      55      60
Gly Ala Phe Gly Lys Val Leu Lys Ser Thr Gln Glu Ala Ile Val Ser
      65      70      75      80
Ser Pro Trp Val Ala Leu Ala Val Arg Ala Arg Pro Gly Val Trp Glu
      85      90      95
His Ile Arg Val Asn Val His Ala Leu Leu Leu Glu His Phe Gln Val
      100      105      110
Asp Glu Tyr Leu His Phe Lys Glu Ala Leu Val Asp Gly Ser Leu Asn
      115      120      125
Pro Asp Ser Glu Pro Leu Thr Ala Thr Phe Gly Arg Arg Pro Phe His
      130      135      140

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```

<210> 48
<211> 90
<212> PRT
<213> Eucalyptus grandis

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```

<400> 48
Gln Glu Ala Ile Val Ser Pro Pro Trp Val Ala Leu Ala Val Arg Pro
 1      5      10      15
Arg Pro Gly Val Trp Glu His Ile Arg Val Asn Val His Ala Leu Val
      20      25      30
Leu Glu Gln Leu Glu Val Ala Glu Tyr Leu His Phe Lys Glu Glu Leu
      35      40      45
Ala Asp Gly Ser Leu Asn Gly Asn Phe Val Leu Glu Leu Asp Phe Glu
      50      55      60
Pro Phe Thr Ala Ser Phe Pro Arg Pro Thr Leu Ser Lys Ser Ile Gly
      65      70      75      80
Asn Gly Val Glu Phe Arg Asn Arg His Leu
      85      90

```

```

<210> 49
<211> 247
<212> PRT

```

## &lt;213&gt; Pinus radiata

&lt;400&gt; 49

```

Met Ala Ala Ala Pro Ala Val Ala Ser Pro Ala Ala Glu Thr Asp Arg
 1          5          10          15
Ile Pro Lys Leu Gln Ala Glu Val Thr Lys Leu Asn Gln Ile Ser Asp
      20          25          30
Asn Glu Lys Glu Gly Phe Val Arg Leu Val Ser Arg Tyr Leu Ser Gly
      35          40          45
Glu Glu Glu Lys Ile Glu Trp Glu Lys Ile Lys Thr Pro Thr Asp Glu
 50          55          60
Ile Val Val Pro Tyr Asp Thr Leu Ala Ala Leu Gly Glu Asp Pro Ser
65          70          75          80
Glu Thr Lys Glu Leu Leu Asp Lys Leu Val Val Leu Lys Leu Asn Gly
      85          90          95
Gly Leu Gly Thr Thr Met Gly Cys Thr Gly Pro Lys Ser Val Ile Glu
      100          105          110
Val Arg Asn Gly Leu Thr Phe Leu Asp Leu Ile Val Lys Gln Ile Glu
      115          120          125
Ser Leu Asn Asn Lys Tyr Asp Ser Lys Val Pro Leu Val Leu Met Asn
130          135          140
Ser Phe Asn Thr His Asp Asp Thr Ile Lys Ile Val Glu Lys Tyr Ser
145          150          155          160
Gly Ser Asn Ile Asp Ile His Ile Phe Asn Gln Ser Gln Tyr Pro Arg
      165          170          175
Met Val Ala Glu Asp Leu Thr Pro Trp Pro Thr Lys Gly Arg Thr Asp
      180          185          190
Lys Glu Ala Trp Tyr Pro Pro Gly His Gly Asp Val Phe Pro Ala Leu
      195          200          205
Leu Asn Ser Gly Lys Leu Asp Glu Leu Leu Ser Gln Gly Lys Glu Tyr
210          215          220
Val Phe Ile Ala Asn Ser Asp Asn Leu Gly Ala Ile Val Asp Leu Ser
225          230          235          240
Ile Leu Phe Ala Leu Val Phe
      245

```

&lt;210&gt; 50

&lt;211&gt; 103

&lt;212&gt; PRT

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 50

```

Met Ala Ala Ala Ala Thr Leu Ser Ala Pro Asp Ala Ala Lys Leu Ser
 1          5          10          15
Gln Leu Lys Ser Ala Val Ser Gly Leu Gly Gln Ile Ser Glu Ser Glu
      20          25          30
Lys Asn Gly Phe Ile Asn Leu Val Ser Arg Tyr Leu Ser Gly Glu Ala
      35          40          45
Gln His Val Asp Trp Ser Lys Ile Gln Thr Pro Thr Asp Glu Ile Val
 50          55          60
Val Pro Tyr Asp Ser Leu Ala Pro Thr Pro Gln Asp Pro Ala Ala Thr
65          70          75          80
Lys Ser Leu Leu Asp Lys Leu Val Val Leu Lys Leu Asn Gly Gly Leu
      85          90          95
Gly Thr Thr Met Gly Cys Thr
      100

```



<210> 51  
 <211> 253  
 <212> PRT  
 <213> Pinus radiata

<400> 51  
 Ala Asn Ser Asp Asn Leu Gly Ala Ile Val Asp Leu Lys Ile Leu Asn  
 1 5 10 15  
 His Leu Val Lys Asn Lys Asn Glu Tyr Cys Met Glu Val Thr Pro Lys  
 20 25 30  
 Thr Leu Ala Asp Val Lys Gly Gly Thr Leu Ile Ser Tyr Glu Gly Arg  
 35 40 45  
 Val Gln Leu Leu Glu Ile Ala Gln Val Pro Glu Glu His Val Gly Glu  
 50 55 60  
 Phe Lys Ser Ile Glu Lys Phe Lys Ile Phe Asn Thr Asn Asn Leu Trp  
 65 70 75 80  
 Val Asn Leu Lys Ala Ile Lys Arg Leu Val Glu Ala Asp Ala Leu Lys  
 85 90 95  
 Met Glu Ile Ile Pro Asn Pro Lys Glu Val Asp Gly Val Lys Val Leu  
 100 105 110  
 Gln Leu Glu Thr Ala Ala Gly Ala Ala Ile Arg Phe Phe Asp Arg Ala  
 115 120 125  
 Ile Gly Val Asn Val Pro Arg Ser Arg Phe Leu Pro Val Lys Ala Thr  
 130 135 140  
 Ser Asp Leu Leu Leu Val Gln Ser Asp Leu Tyr Thr Val Glu Glu Gly  
 145 150 155 160  
 Phe Val Ile Arg Asn Pro Ala Arg Val Asn Pro Thr Asn Pro Thr Ile  
 165 170 175  
 Glu Leu Gly Pro Glu Phe Lys Lys Val Gly Asn Phe Leu Lys Arg Phe  
 180 185 190  
 Lys Ser Ile Pro Ser Ile Ile Asp Leu Asp Ser Leu Lys Val Ser Gly  
 195 200 205  
 Asp Val Trp Phe Gly Ser Gly Val Ile Leu Lys Gly Lys Val Ile Ile  
 210 215 220  
 Glu Ala Lys Gln Gly Ala Thr Leu Glu Ile Pro Asp Glu Ser Val Ile  
 225 230 235 240  
 Glu Asn Lys Val Val Ser Ser Pro Asp Asp Ile Val Asn  
 245 250

<210> 52  
 <211> 184  
 <212> PRT  
 <213> Pinus radiata

<400> 52  
 Met Ser Thr Ile Ile Val Pro Val Pro Ile Pro Thr Pro Ser Glu Asp  
 1 5 10 15  
 Ser Glu Arg Leu Arg Lys Ala Phe Glu Gly Trp Gly Thr Asn Glu Lys  
 20 25 30  
 Ser Ile Ile Gln Ile Leu Gly His Arg Thr Ala Ala Gln Arg Lys Val  
 35 40 45  
 Ile Arg Gln Ser Tyr Phe Gln Leu Tyr Glu Glu Asp Leu Leu Lys Arg  
 50 55 60  
 Leu Glu Ser Glu Leu Ser Ser Asp Phe Glu Lys Ala Val Phe Leu Trp  
 65 70 75 80  
 Val Leu Asp Pro Ala Glu Arg Asp Ala Val Ile Ser His Gly Ala Ile  
 85 90 95

Lys Lys Trp Asn Ala Lys Asn Ile Ser Leu Leu Glu Ile Ser Ser Ala  
 100 105 110  
 Arg Ser Ser Ala Glu Leu Leu Met Val Arg Gln Ala Tyr His Ile Arg  
 115 120 125  
 Asp Lys Lys Ser Leu Glu Glu Asp Val Ala Ala His Thr Ser Gly Asn  
 130 135 140  
 Phe Arg Lys Leu Leu Val Ala Leu Val Ser Ser Tyr Arg Tyr Glu Gly  
 145 150 155 160  
 Pro Glu Val Asp Met His Leu Ala Ser Tyr Glu Ala Lys Lys Leu Ser  
 165 170 175  
 Glu Ser Ile Thr Glu Gln Lys Arg  
 180

<210> 53  
 <211> 213  
 <212> PRT  
 <213> Pinus radiata

<400> 53  
 Met Ala Thr Cys Ser Cys Ala Val Ser Cys Gly Val Asn Pro Val Glu  
 1 5 10 15  
 Arg Asp Cys Glu Glu Ile His Leu Ala Cys Lys Gly Leu Gly Ser Asp  
 20 25 30  
 Glu Glu Lys Ile Ile Glu Ile Leu Gly Ser Lys Asn Glu Gln Gln Arg  
 35 40 45  
 Lys Glu Ile Arg Glu Thr Tyr Tyr Ala Met Tyr Lys Glu Asp Leu Cys  
 50 55 60  
 Lys Arg Leu Glu Lys Glu Leu His Gly Lys Leu Glu Lys Ala Ile Val  
 65 70 75 80  
 Leu Trp Met His Glu Pro Ala Asp Arg Asp Ala Ile Ile Ala Gly Thr  
 85 90 95  
 Ala Leu Glu Gly Trp Cys Thr Asp Asp Arg Ala Leu Ile Glu Val Ile  
 100 105 110  
 Cys Thr Arg Ser Ser Thr Gln Ile Val Lys Ile Arg Glu Ala Tyr Gln  
 115 120 125  
 Lys Arg Tyr Gln Arg Cys Leu Asp Asp Asp Val Ile Cys Lys Thr Asn  
 130 135 140  
 Gly Pro Phe Gln Lys Leu Leu Leu Ala Leu Lys Ala His Arg Cys  
 145 150 155 160  
 Glu Cys Lys Gly Val Asp Ile Asn Lys Ala Arg Cys Asp Ala Lys Met  
 165 170 175  
 Leu Tyr Glu Ala Gly Glu Gly Arg Cys Gly Thr Asp Glu Asp Thr Phe  
 180 185 190  
 Ile Arg Ile Phe Gln Arg Gly Glu Cys Ser Gln Val His Ala Ile Phe  
 195 200 205  
 Ala Cys Asn Lys Gln  
 210

<210> 54  
 <211> 239  
 <212> PRT  
 <213> Eucalyptus grandis

<400> 54  
 Met Ala Thr Ile Ala Val Pro Pro Ser Val Pro Ser Pro Ala Glu Asp  
 1 5 10 15  
 Ala Glu Gln Leu Gln Lys Ala Phe Ala Gly Trp Gly Thr Asn Glu Asp

```

      20      25      30
Leu Ile Ile Ser Ile Leu Pro His Arg Asn Ala Ala Gln Arg Lys Val
      35      40      45
Ile Arg Gln Thr Tyr Ala Glu Thr Tyr Gly Glu Asp Leu Leu Lys Ala
      50      55      60
Leu Asp Lys Glu Leu Ser Ser Asp Phe Glu Arg Ser Val Leu Leu Trp
65      70      75      80
Thr Leu Asp Pro Ala Glu Arg Asp Ala Phe Leu Ser Asn Glu Ala Thr
      85      90      95
Lys Arg Leu Thr Ser Ser Asn Trp Val Leu Met Glu Ile Ala Cys Thr
      100      105      110
Arg Ser Ser Met Glu Leu Phe Met Val Arg Gln Ala Tyr His Ala Arg
      115      120      125
Tyr Lys Lys Ser Leu Glu Glu Asp Ile Ala Tyr His Thr Thr Gly Asp
      130      135      140
Phe Arg Lys Leu Leu Val Pro Leu Ala Ser Thr Phe Arg Tyr Glu Gly
145      150      155      160
Pro Glu Val Asn Met Thr Leu Ala Arg Ser Glu Ala Lys Ile Leu His
      165      170      175
Glu Lys Ile His Glu Lys Ala Tyr Asn His Asp Glu Leu Ile Arg Ile
      180      185      190
Val Thr Thr Arg Ser Lys Ala Gln Leu Asn Ala Thr Leu Asn Tyr Tyr
      195      200      205
Asn Asn Glu Phe Gly Asn Ala Ile Asn Lys Asp Leu Lys Ala Asp Pro
      210      215      220
Asn Asp Glu Phe Leu Lys Leu Leu Arg Ser Ala Ile Lys Cys Leu
225      230      235

```

<210> 55  
 <211> 242  
 <212> PRT  
 <213> Pinus radiata

```

      <400> 55
Met Ser Thr Ile Ile Val Pro Ala Pro Ala Pro Ser Pro Val Glu Asp
1      5      10      15
Ser Glu Arg Leu Arg Lys Ala Phe Glu Gly Trp Gly Thr Asn Glu Lys
      20      25      30
Leu Ile Ile Glu Ile Leu Gly His Arg Thr Ala Ala Gln Arg Arg Ala
      35      40      45
Ile Arg Gln Thr Tyr Thr Gln Leu Tyr Glu Glu Asp Phe Leu Lys Arg
      50      55      60
Leu Gln Ser Glu Leu Thr Arg Asp Phe Glu Arg Ala Leu Leu Leu Trp
65      70      75      80
Ser Leu Asp Pro Pro Glu Arg Asp Ala Leu Leu Ala Tyr Glu Ser Ile
      85      90      95
Lys Lys Trp Ser Pro Asn Asn Arg Ser Leu Leu Glu Ile Ser Ser Ala
      100      105      110
Arg Ser Ser Thr Glu Leu Trp Ser Val Arg Gln Ala Tyr His Ile Arg
      115      120      125
Tyr Lys Lys Ser Leu Glu Glu Asp Val Ala Ser His Thr His Gly Asp
      130      135      140
Phe Arg Lys Leu Leu Val Gln Leu Val Ser Ser Tyr Arg Tyr Glu Gly
145      150      155      160
Pro Glu Val Asp Thr Arg Leu Ala Lys Ser Glu Ala Lys Gln Leu His
      165      170      175
Glu Ala Ile Lys Asp Lys Ala Phe Gly Asn Glu Glu Leu Ile Arg Ile

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			180					185					190				
Ile	Thr	Thr	Arg	Ser	Lys	Ala	Gln	Leu	Asn	Ala	Thr	Phe	Asn	Tyr	Tyr		
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Lys	Asp	Asp	Tyr	Gly	His	His	Ile	Asn	Lys	Asp	Leu	Lys	Glu	Trp	Glu		
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<210> 56  
 <211> 316  
 <212> PRT  
 <213> Eucalyptus grandis

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Leu	Ile	Ile	Ser	Ile	Leu	Gly	His	Arg	Asn	Ala	Ala	Gln	Arg	Lys	Leu		
	35					40						45					
Ile	Arg	Gln	Thr	Tyr	Ala	Glu	Thr	Tyr	Gly	Glu	Asp	Leu	Leu	Lys	Ala		
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Leu	Asp	Arg	Glu	Leu	Thr	Asn	Asp	Phe	Glu	Arg	Leu	Val	Val	Leu	Trp		
65				70					75					80			
Ser	Leu	Asp	Pro	Ala	Glu	Arg	Asp	Ala	Tyr	Leu	Ala	Asn	Glu	Ala	Thr		
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Lys	Arg	Trp	Thr	Ser	Ser	Asn	Gln	Val	Leu	Met	Glu	Ile	Ala	Cys	Thr		
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Arg	Ser	Pro	Gln	Gln	Leu	Leu	Met	Ala	Arg	Gln	Ala	Tyr	His	Ala	Arg		
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Phe	Arg	Lys	Leu	Leu	Val	Pro	Leu	Val	Ser	Ser	Tyr	Arg	Tyr	Asp	Gly		
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			245					250					255				
Arg	Gly	Thr	Asp	Glu	Gly	Ala	Leu	Thr	Arg	Val	Val	Ala	Thr	Arg	Ala		
		260						265				270					
Glu	Val	Asp	Met	Lys	Phe	Ile	Ser	Glu	Glu	Tyr	Gln	Arg	Arg	Asn	Ser		
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Ile	Pro	Leu	Asp	Arg	Ala	Ile	Val	Lys	Asp	Thr	Thr	Gly	Asp	Tyr	Glu		
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&lt;210&gt; 57

&lt;211&gt; 418

&lt;212&gt; DNA

<213> *Eucalyptus grandis*

&lt;400&gt; 57

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&lt;210&gt; 58

&lt;211&gt; 1396

&lt;212&gt; DNA

<213> *Eucalyptus grandis*

&lt;400&gt; 58

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&lt;210&gt; 59

&lt;211&gt; 1861

&lt;212&gt; DNA

<213> *Pinus radiata*

&lt;400&gt; 59

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 <213> Eucalyptus grandis

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<210> 62  
 <211> 460  
 <212> DNA  
 <213> Eucalyptus grandis

<400> 62  
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 <213> Eucalyptus grandis

<400> 63  
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&lt;210&gt; 68

&lt;211&gt; 403

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 68

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&lt;210&gt; 69

&lt;211&gt; 3851

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 69

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&lt;210&gt; 70

&lt;211&gt; 736

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 70

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&lt;210&gt; 71

&lt;211&gt; 448

&lt;212&gt; DNA

## &lt;213&gt; Eucalyptus grandis

## &lt;400&gt; 71

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gaaaaagttt	gatgagaagt	accacttttc	atgtcaattt	actgctgact	tactagccat	360
gaataatgca	gactttatca	tcaccagcac	ataccaagag	atagcaggaa	cgaaaaatac	420
cgttgggcag	tatgagagcc	acactgcc				448

## &lt;210&gt; 72

## &lt;211&gt; 448

## &lt;212&gt; DNA

## &lt;213&gt; Eucalyptus grandis

## &lt;400&gt; 72

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## &lt;210&gt; 73

## &lt;211&gt; 184

## &lt;212&gt; DNA

## &lt;213&gt; Eucalyptus grandis

## &lt;400&gt; 73

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tgat						184

## &lt;210&gt; 74

## &lt;211&gt; 1145

## &lt;212&gt; DNA

## &lt;213&gt; Pinus radiata

## &lt;400&gt; 74

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ctgaa						1145

&lt;210&gt; 75

&lt;211&gt; 1169

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 75

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&lt;210&gt; 76

&lt;211&gt; 420

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 76

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&lt;210&gt; 77

&lt;211&gt; 448

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 77

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 <212> DNA  
 <213> *Eucalyptus grandis*

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 <212> DNA  
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aatggcggtt	tgggaaacaac	catgggttgt	actgggcccc	aatccgtcat	tgaagtacga	660
aatggcctaa	catttctgga	cttgattgtg	aagcaaatag	agtctctgaa	caacaaatac	720
gattccaaag	ttccgttggg	gctgatgaat	tcatttaata	cacatgatga	tacaattaag	780
attgtagaaa	agtactctgg	ctcaaataat	gacatccaca	tcttcaacca	gagtcaatat	840
ccacgcattg	tgccagaaga	tttgacacca	tgcccaacta	aaggctcgta	agacaaagaa	900
gcatggtacc	cacctggcca	cggagatggt	ttccccgctc	ttttgaacag	tggggaagctt	960
gatgaactac	tctcacagg	taaggagtag	gtgttcata	ctaactcaga	caacttggga	1020
gcaatagtgt	atctcaaaat	tttgaatcat	ctgggtcaaaa	ataaaaatga	atactgcatg	1080
gaggtcactc	ccaaaacgct	tgccagatgta	aaagggtggt	cacttatctc	ttatgaaggc	1140
agagttcagc	ttctagagat	tgccacagggt	ccagaggagc	atgttggcga	attcaaggcg	1200
attgaaaagt	tcaaaatctt	caataccaat	aatttatggg	tgaacttgaa	ggcgattaa	1260
aggcttgtgg	aggctgatgc	tcttaaaatg	gagataattc	ctaataccaa	ggaagtggat	1320
ggggtaaagg	tgcttcagct	tgaaacggct	gcagggtgct	ctataagggt	ttttgatcgt	1380
gcaattgggt	tcaatgttcc	aagatcacgc	tttctcccag	tgaaggcaac	ttcagattta	1440
ctacttgtgc	agtcggatct	ttacactgtg	gaggaagggt	ttgtcatccg	aaatcctgct	1500
agagtcaacc	ctacaaatcc	caccattgag	ttgggtcctg	aattcaagaa	ggttggcaac	1560
tttctaaagc	gtttcaagtc	catacctagc	atcattgatc	ttgatagtct	aaagggtgtca	1620
gggtgatgtg	ggtttggcag	tgaggtcatc	ctgaaggga	aagttataat	tgaagcaag	1680
cagggggcta	cacttgagat	acctgatgaa	tctgtgatag	aaaacaagg	agtgagcagt	1740
cctgatgata	ttgtgaacta	gcttataagt	tttggcagtt	ggcattcttg	atttttgggt	1800
tggtttaga	gctagaatac	tgtgtttagt	gcccttaaag	agtaggggtg	aagtttcacc	1860
tctgggttta	tcttccataa	tccttactgg	gaggggtttc	ataattttat	tataactagc	1920
cagtttctat	gtaaatgttg	agcgcaatt	tttatgcatg	aaagcctagg	atgaagcatc	1980
caaacacaat	aaataaatgg	tgctttgttg	agatttttta	aaaaaaaaa	aaaaaaaaa	2040
aaaaa						2045

<210> 81  
 <211> 139  
 <212> PRT  
 <213> Eucalyptus grandis

&lt;400&gt; 81

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Phe Asp Phe Ala Arg Gly Tyr Ser Pro Lys Tyr Val Lys Glu Tyr Ile
 1          5          10          15
Glu Ser Ala Lys Pro Leu Phe Ser Val Gly Glu Tyr Trp Asp Ser Cys
          20          25          30
Asn Tyr Ser Gly Thr Thr Leu Glu Tyr Asn Gln Asp Ser His Arg Gln
          35          40          45
Arg Ile Val Asn Trp Ile Asp Gly Thr Gly Gln Leu Ser Ala Ala Phe
          50          55          60
Asp Phe Thr Thr Lys Gly Ile Leu Gln Glu Ala Val Lys Gly Gln Phe
          65          70          75          80
Trp Arg Leu Arg Asp Pro Lys Gly Lys Pro Pro Gly Val Met Gly Trp
          85          90          95
Trp Pro Ser Arg Ala Val Thr Phe Leu Asp Asn His Asp Thr Gly Ser
          100          105          110
Thr Gln Ala His Trp Pro Phe Pro Ser Asn His Ile Arg Arg Val Thr
          115          120          125
Arg Thr Tyr Ser Leu Ile Gln Glu Tyr Leu Leu
          130          135

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&lt;210&gt; 82

&lt;211&gt; 189

&lt;212&gt; PRT

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 82

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Leu Phe Trp Leu Phe Val Leu Leu Val Phe Tyr Leu Ala Ala Ser Ala
 1          5          10          15
Ser Pro Ala Leu Leu Phe Gln Gly Phe Asn Trp Glu Ser Trp Lys Lys
          20          25          30
Glu Gly Gly Trp Tyr Asn Ser Leu Lys Asn Leu Val Pro Asp Leu Ala
          35          40          45
Asn Ala Gly Ile Thr His Val Trp Leu Pro Pro Pro Ser Gln Ser Ala
          50          55          60
Ala Gln Gln Gly Tyr Leu Pro Gly Arg Leu Tyr Asp Leu Asn Ala Ser
          65          70          75          80
Ser Tyr Gly Asn Gln Asp Glu Leu Lys His Leu Ile Asp Ala Phe His
          85          90          95
Gln Lys Gly Ile Lys Cys Leu Ala Asp Ile Val Ile Asn His Arg Thr
          100          105          110
Ala Glu Lys Gln Asp Ser Arg Gly Ile Trp Cys Ile Phe Glu Gly Gly
          115          120          125
Thr Pro Asp Asp Arg Leu Asp Trp Gly Pro Ser Leu Ile Cys Arg Asp
          130          135          140
Asp Thr Glu Tyr Ser Asp Gly Arg Gly Asn Leu Asp Ser Gly Glu Asp
          145          150          155          160
Phe Lys Pro Ala Pro Asp Ile Asp His Leu Asn Pro Arg Val Gln Lys
          165          170          175
Glu Leu Ser Asp Trp Met Asn Trp Leu Lys Ser Asp Ile
          180          185

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&lt;210&gt; 83

&lt;211&gt; 176

&lt;212&gt; PRT

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 83

Phe Gly Glu Phe Asn Thr Asp Glu Met Ala Gly Val Met Ala Ala Gly  
 1 5 10 15  
 Val Ala Asn Leu Asn Val Leu Gly Arg Glu Thr Ala Glu Phe Thr Ser  
 20 25 30  
 Phe Arg Pro Val Phe Leu Arg Gly Asn Ser Gln Gly Leu Ser Ser Ala  
 35 40 45  
 Ser Ser Leu Cys Asp Tyr Arg Ile Phe Ala Asp Ser Lys Arg Lys Lys  
 50 55 60  
 His Ala Ile Phe Arg Lys Gln Asn Ile Asn Arg Ser Thr Val Val Ser  
 65 70 75 80  
 Pro Arg Ala Val Ser Asp Thr Phe Ser Glu Leu Thr Cys Leu Asp Pro  
 85 90 95  
 Val Ala Ser Arg Ser Val Leu Gly Ile Ile Leu Gly Gly Gly Ala Gly  
 100 105 110  
 Thr Arg Leu Tyr Pro Leu Thr Lys Lys Arg Ala Lys Pro Ala Val Pro  
 115 120 125  
 Leu Gly Ala Asn Tyr Arg Leu Ile Asp Ile Pro Val Ser Asn Cys Ile  
 130 135 140  
 Asn Ser Asn Ile Ser Lys Ile Tyr Val Leu Thr Gln Phe Asn Ser Ala  
 145 150 155 160  
 Ser Leu Asn Pro Ser Ser Phe Thr Gly Leu Phe Lys His Met Gly Ser  
 165 170 175

&lt;210&gt; 84

&lt;211&gt; 47

&lt;212&gt; PRT

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 84

Asp Pro Ala Leu Asp Ser Ala Asp Ala Phe Lys Ser Val Arg Arg Asp  
 1 5 10 15  
 Pro Asp Val Val Ser Pro Arg Asp Val Ser Asp Ser Arg Asn Ser Gln  
 20 25 30  
 Thr Cys Leu Asn Pro Asp Ala Ser Arg Ser Val Leu Gly Ile Ile  
 35 40 45

&lt;210&gt; 85

&lt;211&gt; 146

&lt;212&gt; PRT

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 85

Ala Pro Ala Leu Ala Ser Gly Ala Ala Ala Phe Lys Ser Val Arg Arg  
 1 5 10 15  
 Ala Pro Ala Val Val Ser Pro Arg Ala Val Ser Asp Ser Arg Asn Ser  
 20 25 30  
 Gln Thr Cys Leu Asp Pro Asp Ala Ser Arg Ser Val Leu Gly Ile Ile  
 35 40 45  
 Leu Gly Gly Gly Ala Gly Thr Arg Leu Tyr Pro Leu Thr Lys Lys Arg  
 50 55 60  
 Ala Lys Pro Ala Val Pro Leu Gly Ala Asn Tyr Arg Leu Ile Asp Ile  
 65 70 75 80  
 Pro Val Ser Asn Cys Leu Asn Ser Asn Val Ser Lys Ile Tyr Val Leu  
 85 90 95  
 Thr Gln Phe Asn Ser Ala Ser Leu Asn Arg His Leu Ser Arg Ala Tyr  
 100 105 110



Ala Ser Asn Met Gly Gly Tyr Lys Asn Glu Gly Phe Val Glu Val Leu  
 115 120 125  
 Ala Ala Gln Gln Ser Pro Glu Asn Pro Asn Trp Phe Gln Gly Thr Ala  
 130 135 140  
 Asp Ala  
 145

<210> 86  
 <211> 84  
 <212> PRT  
 <213> Eucalyptus grandis

<400> 86  
 Glu Leu Thr Ala Met Asp Ser Arg Cys Val Ala Leu Lys Ala Asn Ala  
 1 5 10 15  
 Ser Leu Ala Gln Ser Asn Lys Ser Cys Leu Lys Asn Val Asp Lys Gly  
 20 25 30  
 Phe Leu Gly Glu Arg Ile Arg Gly Ser Leu Asp Asn Ser Val Trp Val  
 35 40 45  
 Lys Gln Val Ala Arg Asn Leu Arg Val Glu Lys Lys Phe Lys Lys Ala  
 50 55 60  
 Lys Pro Gly Val Ala Phe Ala Val Ile Thr Ser Asn Thr Val Ala Glu  
 65 70 75 80  
 Thr Leu Thr Ile

<210> 87  
 <211> 113  
 <212> PRT  
 <213> Eucalyptus grandis

<400> 87  
 Lys Ile Phe Ile Leu Thr Gln Phe Asn Ser Phe Ser Leu Asn Arg His  
 1 5 10 15  
 Leu Ser Arg Thr Tyr Asn Phe Asp Asn Gly Val Ser Phe Gly Asp Gly  
 20 25 30  
 Phe Val Glu Val Leu Ala Ala Thr Gln Thr Pro Gly Glu Ala Gly Lys  
 35 40 45  
 Arg Trp Phe Gln Gly Thr Ala Asp Ala Val Arg Gln Phe Ile Trp Val  
 50 55 60  
 Phe Glu Asp Ala Lys Asn Lys Asn Val Glu Asn Ile Leu Ile Leu Ser  
 65 70 75 80  
 Gly Asp His Leu Tyr Arg Met Asn Tyr Met Asp Phe Val Gln Lys His  
 85 90 95  
 Ile Asp Ser Asn Ala Asp Ile Thr Val Ser Cys Val Pro Met Asp Asp  
 100 105 110  
 Ser

<210> 88  
 <211> 131  
 <212> PRT  
 <213> Eucalyptus grandis

<400> 88  
 Ala Ile Ser Val Ala Cys Thr Pro Val Gly Glu Ser Arg Ala Ser Asp  
 1 5 10 15

Tyr Gly Leu Val Lys Ile Asp Ser Arg Gly Gln Ile Val Gln Phe Ser  
                   20                  25                  30  
 Glu Lys Pro Lys Gly Pro Asp Leu Thr Ala Met Gln Val Asp Thr Thr  
                   35                  40                  45  
 Thr Leu Gly Leu Ser Pro Gln Glu Ala Ala Arg Ser Pro Tyr Ile Ala  
                   50                  55                  60  
 Ser Met Gly Val Tyr Ala Phe Lys Thr Glu Ser Leu Leu Asn Leu Leu  
   65                  70                  75                  80  
 Lys Trp Arg Tyr Pro Thr Ala Asn Asp Phe Gly Ser Glu Ile Ile Pro  
                   85                  90                  95  
 Ser Ala Val Met Glu Gln Asp Val Gln Ala Tyr Ile Phe Arg Asp Tyr  
                   100                  105                  110  
 Trp Glu Asp Ile Gly Thr Ile Lys Ser Phe Tyr Asp Ala Asn Leu Ala  
                   115                  120                  125  
 Leu Thr Glu  
                   130

<210> 89  
 <211> 115  
 <212> PRT  
 <213> Eucalyptus grandis

<400> 89  
 Arg Leu Ser Ser Lys Phe Ile Trp Val Trp Leu Leu Leu Arg Trp Val  
   1                  5                  10                  15  
 Arg Phe Val Gly Met Asp Ser Cys Phe Ala Ser Met Lys Val Gly Ala  
                   20                  25                  30  
 Arg Pro Val Pro Gly Gly Gly Ile Ile Asn Phe Ser Glu Phe Trp Gly  
                   35                  40                  45  
 Glu Asn Leu Arg Val Gly Ala Asn Lys Gln Phe Gly Ala Arg Leu Cys  
   50                  55                  60  
 Lys Ser Leu Arg Ser Glu Thr Arg Ile Gly Arg Val Lys Pro Gly Ile  
   65                  70                  75                  80  
 Ala Tyr Ser Val Leu Thr Pro Glu Val Asp Lys Glu Thr Met Thr Leu  
                   85                  90                  95  
 Gln Ala Pro Val Leu Glu Thr Pro Arg Ala Asp Pro Lys Ser Phe Ala  
                   100                  105                  110  
 Ser Ile Ile  
                   115

<210> 90  
 <211> 600  
 <212> PRT  
 <213> Eucalyptus grandis

<400> 90  
 Pro Gly Met Ile Gln Val Tyr Leu Gly Ser Ala Gly Ala Leu Asp Val  
   1                  5                  10                  15  
 Glu Gly Lys Glu Leu Pro Arg Leu Val Tyr Val Ser Arg Glu Lys Arg  
                   20                  25                  30  
 Pro Gly Tyr Gln His His Lys Lys Ala Gly Ala Met Asn Ala Leu Val  
                   35                  40                  45  
 Arg Val Ser Ala Val Leu Thr Asn Ala Pro Phe Leu Leu Asn Leu Asp  
   50                  55                  60  
 Cys Asp His Tyr Ile Asn Asn Ser Lys Ala Ile Arg Glu Ala Met Cys  
   65                  70                  75                  80  
 Phe Leu Met Asp Pro Gln Leu Gly Lys Lys Leu Cys Tyr Val Gln Phe

															85										90										95				
Pro	Gln	Arg	Phe	Asp	Gly	Ile	Asp	Arg	His	Asp	Arg	Tyr	Ala	Asn	Arg																								
				100					105								110																						
Asn	Ile	Val	Phe	Phe	Asp	Ile	Asn	Met	Arg	Gly	Leu	Asp	Gly	Ile	Gln																								
				115					120								125																						
Gly	Pro	Val	Tyr	Val	Gly	Thr	Gly	Cys	Val	Phe	Asn	Arg	Gln	Ala	Leu																								
				130					135								140																						
Tyr	Gly	Tyr	Asp	Pro	Pro	Val	Ser	Gln	Lys	Arg	Pro	Lys	Met	Thr	Cys																								
				145					150								155																						
Asp	Cys	Trp	Pro	Ser	Trp	Cys	Ser	Cys	Cys	Cys	Gly	Gly	Ser	Arg	Lys																								
				165					170								175																						
Ser	Lys	Ser	Lys	Lys	Lys	Asp	Asp	Thr	Ser	Leu	Leu	Gly	Pro	Val	His																								
				180					185								190																						
Ala	Lys	Lys	Lys	Lys	Met	Thr	Gly	Lys	Asn	Tyr	Leu	Lys	Lys	Lys	Gly																								
				195					200								205																						
Ser	Gly	Pro	Val	Phe	Asp	Leu	Glu	Asp	Ile	Glu	Glu	Gly	Leu	Glu	Gly																								
				210					215								220																						
Phe	Asp	Glu	Leu	Glu	Lys	Ser	Ser	Leu	Met	Ser	Gln	Lys	Asn	Phe	Glu																								
				225					230								235																						
Lys	Arg	Phe	Gly	Gln	Ser	Pro	Val	Phe	Ile	Ala	Ser	Thr	Leu	Met	Glu																								
				245					250								255																						
Asp	Gly	Gly	Leu	Pro	Glu	Gly	Thr	Asn	Ser	Thr	Ser	Leu	Ile	Lys	Glu																								
				260					265								270																						
Ala	Ile	His	Val	Ile	Ser	Cys	Gly	Tyr	Glu	Glu	Lys	Thr	Glu	Trp	Gly																								
				275					280								285																						
Lys	Glu	Ile	Gly	Trp	Ile	Tyr	Gly	Ser	Val	Thr	Glu	Asp	Ile	Leu	Thr																								
				290					295								300																						
Gly	Phe	Lys	Met	His	Cys	Arg	Gly	Trp	Lys	Ser	Val	Tyr	Cys	Met	Pro																								
				305					310								315																						
Lys	Arg	Pro	Ala	Phe	Lys	Gly	Ser	Ala	Pro	Ile	Asn	Leu	Ser	Asp	Arg																								
				325					330								335																						
Leu	His	Gln	Val	Leu	Arg	Trp	Ala	Leu	Gly	Ser	Val	Glu	Ile	Phe	Leu																								
				340					345								350																						
Ser	Arg	His	Cys	Pro	Leu	Trp	Tyr	Ala	Trp	Gly	Gly	Lys	Leu	Lys	Leu																								
				355					360								365																						
Leu	Glu	Arg	Leu	Ala	Tyr	Ile	Asn	Thr	Ile	Val	Tyr	Pro	Phe	Thr	Ser																								
				370					375								380																						
Ile	Pro	Leu	Leu	Phe	Tyr	Cys	Thr	Ile	Pro	Ala	Val	Cys	Leu	Leu	Thr																								
				385					390								395																						
Gly	Lys	Phe	Ile	Ile	Pro	Thr	Leu	Thr	Asn	Phe	Ala	Ser	Ile	Trp	Phe																								
				405					410								415																						
Leu	Ala	Leu	Phe	Leu	Ser	Ile	Ile	Ala	Thr	Gly	Val	Leu	Glu	Leu	Arg																								
				420					425								430																						
Trp	Ser	Gly	Val	Ser	Ile	Glu	Asp	Trp	Trp	Arg	Asn	Glu	Gln	Phe	Trp																								
				435					440								445																						
Val	Ile	Gly	Gly	Val	Ser	Ala	His	Leu																															

Leu Tyr Pro Phe Leu Lys Gly Leu Met Gly Lys Gln Asn Arg Thr Pro  
 545 550 555 560  
 Thr Ile Val Val Leu Trp Ser Val Leu Leu Ala Ser Ile Phe Ser Leu  
 565 570 575  
 Val Trp Val Arg Ile Asp Pro Phe Leu Pro Lys Gln Thr Gly Pro Val  
 580 585 590  
 Leu Lys Pro Cys Gly Val Glu Cys  
 595 600

&lt;210&gt; 91

&lt;211&gt; 222

&lt;212&gt; PRT

&lt;213&gt; Pinus radiata

&lt;400&gt; 91

Pro Asn Glu Phe Pro Leu Tyr Thr Thr Leu Glu Lys Lys Ser Leu Leu  
 1 5 10 15  
 Tyr Arg Ala Tyr Ser Cys Thr His Phe Cys Ala Ile Ile Gly Leu Ile  
 20 25 30  
 Cys Tyr Arg Leu Leu Tyr Ile Pro Ser Glu Asp Ser Trp Ser Trp Ile  
 35 40 45  
 Leu Ile Phe Val Ala Glu Leu Gly Phe Ser Tyr Ser Trp Ile Leu Asp  
 50 55 60  
 Gln Ala Leu Arg Trp Trp Pro Val Gln Arg Thr Val Phe Pro Lys Arg  
 65 70 75 80  
 Leu Ser Lys Arg Phe Gln Ser Asn Leu Pro Pro Val Asp Ile Phe Ile  
 85 90 95  
 Cys Thr Ala Asp Pro Phe Lys Glu Pro Pro Leu Thr Val Ile Asn Thr  
 100 105 110  
 Val Leu Ser Ala Leu Ala Val His Tyr Pro Met Gly Lys Leu Ser Cys  
 115 120 125  
 Tyr Val Ser Asp Asp Gly Gly Ser Pro Leu Thr Phe Tyr Ala Leu Leu  
 130 135 140  
 Glu Ala Ser Arg Phe Ala Lys Ile Trp Ile Pro Phe Cys Asp Lys Tyr  
 145 150 155 160  
 Ser Ile Glu Asp Arg Cys Pro Glu Val Tyr Phe Ser Asn Pro Ser Ala  
 165 170 175  
 Leu Glu Asn Val Asn Leu Ser Phe Met Thr Asp Trp Arg His Val Asn  
 180 185 190  
 Lys Met Tyr Phe Glu Leu Lys Asp Arg Ile Asn Asn Val Met Glu Met  
 195 200 205  
 Gly Ser Val His Gln Ile Asn Arg Met Asn Thr Lys Asp Ser  
 210 215 220

&lt;210&gt; 92

&lt;211&gt; 121

&lt;212&gt; PRT

&lt;213&gt; Pinus radiata

&lt;400&gt; 92

Ser Lys Leu Leu Met Glu Pro Asn Asp Phe Pro Leu Tyr Thr Thr Leu  
 1 5 10 15  
 Glu Lys Lys Ser Leu Leu Tyr Arg Ala Tyr Ser Cys Thr His Phe Ser  
 20 25 30  
 Ala Ile Ile Gly Leu Ile Cys Tyr Arg Leu Leu Tyr Ile Pro Ser Glu  
 35 40 45  
 Asp Ser Trp Pro Trp Ile Leu Ile Phe Val Ala Glu Leu Gly Phe Ser

50                      55                      60  
 Tyr Ser Trp Ile Leu Asp Gln Ala Leu Arg Trp Trp Pro Val Glu Arg  
 65                      70                      75                      80  
 Thr Val Phe Pro Asn Arg Leu Ser Lys Arg Phe Gln Ser Lys Leu Pro  
                     85                      90                      95  
 Pro Val Asp Ile Phe Ile Cys Thr Ala Asp Pro Phe Lys Glu Pro Pro  
                     100                      105                      110  
 Leu Thr Val Ile Asn Thr Val Leu Ser  
                     115                      120

<210> 93  
 <211> 603  
 <212> PRT  
 <213> Pinus radiata

<400> 93  
 Leu Lys Phe Phe Glu Phe Gly Arg Glu Leu Asn Leu Thr Met Glu Ala  
 1                      5                      10                      15  
 Ser Ala Gly Leu Val Ala Gly Ser His Asn Arg Asn Glu Phe Val Val  
                     20                      25                      30  
 Ile His Gly His Glu Glu Pro Lys Pro Leu Asn Thr Leu Ser Gly His  
                     35                      40                      45  
 Val Cys Gln Ile Cys Gly Glu Asp Val Gly Leu Asn Thr Asp Gly Glu  
                     50                      55                      60  
 Leu Phe Val Ala Cys Asn Glu Cys Gly Phe Pro Val Cys Arg Pro Cys  
 65                      70                      75                      80  
 Tyr Glu Tyr Glu Arg Arg Glu Gly Asn Gln Ser Cys Pro Gln Cys Asn  
                     85                      90                      95  
 Thr Arg Tyr Lys Arg Gln Lys Gly Ser Pro Arg Val Glu Gly Asp Asp  
                     100                      105                      110  
 Asp Glu Glu Asp Val Asp Asp Ile Glu His Glu Phe Asn Val Glu Thr  
                     115                      120                      125  
 Gln Gln Arg Asn Arg Gln Gln Ile Thr Glu Ala Met Leu His Gly Arg  
                     130                      135                      140  
 Met Ser Tyr Gly Arg Gly Pro Asp Asp Glu Asn Ser Gln Ile Ala His  
 145                      150                      155                      160  
 Asn Pro Glu Leu Pro Pro Gln Ile Pro Val Leu Ala Asn Gly His Ser  
                     165                      170                      175  
 Val Val Ser Gly Glu Ile Pro Thr Ser Tyr Tyr Ala Asp Asn Gln Leu  
                     180                      185                      190  
 Leu Ala Asn Pro Ala Met Leu Lys Arg Val His Pro Ser Ser Glu Pro  
                     195                      200                      205  
 Gly Ser Gly Arg Ile Ile Met Asp Pro Asn Arg Asp Ile Gly Ser Tyr  
                     210                      215                      220  
 Gly Phe Gly Asn Val Ser Trp Lys Glu Arg Gly Asp Gly Tyr Lys Ser  
 225                      230                      235                      240  
 Lys Glu Asn Lys Ser Gly Gln Leu Asp Met Thr Glu Gly Arg Tyr Gln  
                     245                      250                      255  
 Tyr Asn Gly Gly Phe Ala Pro Asn Glu Pro Glu Asp Tyr Ile Asp Pro  
                     260                      265                      270  
 Asp Met Pro Met Thr Asp Glu Ala Arg Gln Pro Leu Ser Arg Lys Val  
                     275                      280                      285  
 Pro Ile Pro Ser Ser Lys Ile Asn Pro Tyr Arg Met Val Ile Val Ile  
                     290                      295                      300  
 Arg Leu Ile Val Leu Gly Ile Phe Leu Arg Tyr Arg Leu Leu Asn Pro  
 305                      310                      315                      320  
 Val Lys Asn Ala Tyr Gly Leu Trp Ala Thr Ser Ile Val Cys Glu Ile

325 330 335  
 Trp Phe Ala Leu Ser Trp Ile Leu Asp Gln Phe Pro Lys Trp Leu Pro  
 340 345 350  
 Ile Ser Arg Glu Thr Tyr Leu Asp Arg Leu Ser Leu Arg Tyr Glu Arg  
 355 360 365  
 Glu Gly Glu Pro Ser Met Leu Ala Pro Val Asp Leu Phe Val Ser Thr  
 370 375 380  
 Val Asp Pro Leu Lys Glu Pro Pro Leu Val Thr Ala Asn Thr Val Leu  
 385 390 395 400  
 Ser Ile Leu Ser Val Asp Tyr Pro Val Asp Asn Val Ser Cys Tyr Val  
 405 410 415  
 Ser Asp Asp Gly Ala Ser Met Leu Thr Phe Glu Ser Leu Ser Glu Thr  
 420 425 430  
 Ser Glu Phe Ala Arg Lys Trp Val Pro Phe Cys Lys Lys Phe Asp Ile  
 435 440 445  
 Glu Pro Arg Ala Pro Glu Ile Tyr Phe Ser Gln Lys Ile Asp Tyr Leu  
 450 455 460  
 Lys Asp Lys Phe Gln Pro Thr Phe Val Lys Glu Arg Arg Ala Met Lys  
 465 470 475 480  
 Arg Glu Tyr Glu Glu Phe Lys Val Arg Ile Asn Arg Leu Val Ala Lys  
 485 490 495  
 Ala Ser Lys Val Pro Lys Glu Gly Trp Thr Met Gln Asp Gly Thr Pro  
 500 505 510  
 Trp Pro Gly Asn Asn Thr Arg Asp His Pro Gly Met Ile Gln Val Phe  
 515 520 525  
 Leu Gly His Ser Gly Gly Leu Asp Thr Glu Gly Asn Glu Leu Pro Arg  
 530 535 540  
 Leu Val Tyr Val Ser Arg Glu Lys Arg Pro Gly Phe Gln His His Lys  
 545 550 555 560  
 Lys Ala Gly Ala Met Asn Ala Leu Val Arg Val Ser Ala Val Leu Thr  
 565 570 575  
 Asn Ala Pro Phe Met Leu Asn Leu Asp Cys Asp His Tyr Ile Asn Asn  
 580 585 590  
 Ser Lys Ala Ile Arg Glu Gly Met Cys Phe Met  
 595 600

&lt;210&gt; 94

&lt;211&gt; 245

&lt;212&gt; PRT

&lt;213&gt; Pinus radiata

&lt;400&gt; 94

Asn His Ile Lys Leu Leu Pro Phe Ala Gln Glu Gln Asn Asp Glu Ile  
 1 5 10 15  
 Met Glu Ala Arg Ala Gly Leu Val Ala Gly Ser Tyr Lys Arg Asn Glu  
 20 25 30  
 Leu Met Val Val Pro Gly His Asp Gly Pro Lys Pro Ile Arg Leu Ser  
 35 40 45  
 Thr Leu Gln Asp Cys Gln Val Cys Gly Asp Lys Ile Gly Cys Asn Pro  
 50 55 60  
 Asn Gly Glu Leu Phe Val Ala Cys Asn Glu Cys Gly Phe Pro Val Cys  
 65 70 75 80  
 Arg Pro Cys Tyr Glu Tyr Glu Arg Lys Asp Gly Asn Arg Cys Cys Pro  
 85 90 95  
 Gln Cys Lys Thr Arg Tyr Arg Arg His Lys Gly Ser Pro Arg Val Glu  
 100 105 110  
 Gly Asp Asp Glu Glu Asp Gly Met Asp Asp Leu Glu Gln Glu Phe Asn

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      115      120      125
Met Glu Arg Asp Arg Gln Ser Val Val Ser His Arg Gly Asn Ala Phe
  130      135      140
Asp Ala Thr Pro Arg Ala Ala His Ser Ile Ala Asn Arg Ser Ile Asn
145      150      155      160
Gly Asp Asn Tyr Ala Leu Ser Leu Pro Pro Ile Met Asp Gly Asp Ser
      165      170      175
Leu Ser Val Gln Arg Phe Pro His Ala Ala Thr Val Ile Gly Asn Gly
      180      185      190
Leu Asp Pro Val Lys Glu Asn Tyr Gly Ser Ala Ala Trp Lys Glu Arg
  195      200      205
Val Glu Asn Trp Lys Ala Lys His Asp Lys Lys Ser Gly Ser Ile Lys
  210      215      220
Asp Gly Ile Tyr Asp Pro Asp Glu Ala Asp Asp Ile Met Met Thr Glu
225      230      235      240
Ala Glu Ala Arg Gln
      245

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<210> 95
<211> 149
<212> PRT
<213> Eucalyptus grandis

```

```

      <400> 95
Leu Glu Arg Val Ser Gly Thr Glu His Ser His Ile Leu Arg Val Pro
  1      5      10      15
Phe Arg Ser Asp Gln Gly Ile Leu Arg Lys Trp Ile Ser Arg Phe Asp
      20      25      30
Val Trp Pro Tyr Leu Glu Thr Phe Ala Leu Asp Ala Ala His Glu Ile
      35      40      45
Thr Ala Glu Leu Gln Gly Phe Pro Asp Phe Ile Ile Gly Asn Tyr Ser
  50      55      60
Asp Gly Asn Leu Val Ala Ser Leu Leu Ala Tyr Lys Met Gly Val Thr
65      70      75      80
Gln Cys Thr Ile Ala His Ala Leu Glu Lys Thr Lys Tyr Pro Asp Ser
      85      90      95
Asp Ile Tyr Trp Lys Lys Phe Asp Glu Lys Tyr His Phe Ser Cys Gln
      100      105      110
Phe Thr Ala Asp Leu Leu Ala Met Asn Asn Ala Asp Phe Ile Ile Thr
      115      120      125
Ser Thr Tyr Gln Glu Ile Ala Gly Thr Lys Asn Thr Val Gly Gln Tyr
  130      135      140
Glu Ser His Thr Ala
145

```

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<210> 96
<211> 124
<212> PRT
<213> Eucalyptus grandis

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      <400> 96
Leu Ala Lys Ala Gly Thr Lys Asn Thr Val Gly Gln Tyr Glu Ser His
  1      5      10      15
Thr Ala Phe Thr Leu Pro Gly Leu Tyr Arg Val Val His Gly Ile Asp
      20      25      30
Val Phe Asp Pro Lys Phe Asn Ile Val Ser Pro Gly Ala Asp Met Cys
      35      40      45

```

Ile Tyr Phe Pro Tyr Ser Glu Lys Gln Lys Arg Leu Thr Ala Leu His  
 50 55 60  
 Gly Ser Ile Glu Lys Leu Leu Tyr Asp Pro Glu Gln Asn Asp Glu His  
 65 70 75 80  
 Ile Gly Ser Leu Ser Asp Arg Ser Lys Pro Met Ile Phe Ser Met Ala  
 85 90 95  
 Arg Leu Asp Lys Val Lys Asn Met Thr Gly Leu Val Glu Cys Tyr Ala  
 100 105 110  
 Lys Asn Ser Lys Leu Arg Glu Leu Ala Asn Leu Val  
 115 120

<210> 97  
 <211> 61  
 <212> PRT  
 <213> Eucalyptus grandis

<400> 97  
 Glu Ala Ile Val Leu Pro Pro Phe Val Ala Ile Ala Val Arg Pro Arg  
 1 5 10 15  
 Pro Gly Val Trp Glu Tyr Val Arg Val Asn Val His Glu Leu Ser Val  
 20 25 30  
 Glu Gln Leu Thr Val Ser Glu Tyr Leu Gly Phe Lys Glu Glu Leu Val  
 35 40 45  
 Asp Gly Lys Ser Glu Asp Ser Phe Val Leu Glu Leu Asp  
 50 55 60

<210> 98  
 <211> 217  
 <212> PRT  
 <213> Pinus radiata

<400> 98  
 Cys Val Gly Ile Asp Pro Lys Ala Asn Met Val Ser Ala Arg Leu Thr  
 1 5 10 15  
 Arg Ser Leu Ser Ser Arg Glu Arg Val Glu Asp Thr Leu Ser Glu His  
 20 25 30  
 Arg Asn Gln Leu Ala Ala Leu Phe Ser Arg Tyr Val Ala Gln Gly Lys  
 35 40 45  
 Lys Val Leu Gln Pro His Glu Ile Leu Asp Gly Leu Ala Ala Val Ile  
 50 55 60  
 Gly Glu Asn Asp Glu His Gln Asn Phe Arg Asp Gly Leu Phe Gly Asn  
 65 70 75 80  
 Val Leu Arg Ser Thr Gln Glu Ala Ile Ile Ile Pro Pro Trp Val Val  
 85 90 95  
 Leu Ala Val Arg Pro Arg Pro Gly Val Trp Glu Phe Val Arg Val Asn  
 100 105 110  
 Val Asp Glu Leu Ala Val Glu Gln Leu Ser Val Ala Glu Tyr Leu Glu  
 115 120 125  
 Phe Lys Glu Gln Leu Val Asp Gly Ser Val Lys Asp Asn Tyr Val Leu  
 130 135 140  
 Glu Leu Asp Leu Glu Pro Phe Asn Ala Ser Phe Pro Arg Pro Thr Gln  
 145 150 155 160  
 Pro Ser Ser Ile Gly Ser Gly Val Gln Phe Leu Asn Arg His Leu Ser  
 165 170 175  
 Ser Arg Leu Phe Arg Asp His Glu Ser Met Gln Pro Leu Leu Asp Phe  
 180 185 190  
 Leu Arg Ala His Lys Tyr Gln Gly Gln Arg Leu Met Leu Asn Glu Arg



[illegible]

&lt;212&gt; PRT

&lt;213&gt; Pinus radiata

&lt;400&gt; 100

```

Ser Asn Leu Glu Thr Phe Leu Gly Arg Val Pro Met Val Phe Asn Val
 1          5          10          15
Val Ile Leu Ser Pro His Gly Tyr Phe Gly Gln Ala Asn Val Leu Gly
          20          25          30
Met Pro Asp Thr Gly Gly Gln Val Val Tyr Ile Leu Asp Gln Cys Arg
          35          40          45
Ala Leu Glu Asn Glu Met Leu Leu Arg Ile Lys Gln Gln Gly Leu Asp
          50          55          60
Ile Thr Pro Glu Ile Ile Val Val Thr Arg Leu Ile Pro Glu Ala His
65          70          75          80
Gly Thr Thr Cys Asn Gln Arg Leu Glu Lys Ile Ser Gly Thr Gln His
          85          90          95
Ser Arg Ile Leu Arg Val Pro Phe Arg Thr Glu Lys Gly Val Val Arg
          100          105          110
Asp Trp Val Ser Arg Phe Asp Val Trp Pro Tyr Leu Glu Arg Phe Ser
          115          120          125
Glu Asp Val Thr Asn Glu Ile Ala Val Glu Leu
130          135

```

&lt;210&gt; 101

&lt;211&gt; 68

&lt;212&gt; PRT

&lt;213&gt; Pinus radiata

&lt;400&gt; 101

```

Ile Leu Leu Leu Ile Val Gly Ile Gly Ile His Ile Lys Ala Lys Glu
 1          5          10          15
Asn Met Val Ala Ala Ala Leu Thr His Ala Leu Ser Ser Arg Glu Arg
          20          25          30
Val Glu Asp Met Leu Ser Glu His Arg Asn Glu Ile Val Ser Leu Leu
          35          40          45
Ser Arg Tyr Val Ala Glu Gly Lys Lys Ile Leu Gln Pro His Gln Leu
          50          55          60
Leu Asp Gly Leu
65

```

&lt;210&gt; 102

&lt;211&gt; 70

&lt;212&gt; PRT

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 102

```

Met Ala Ala Pro Lys Leu Gly Arg Ile Pro Ser Ile Arg Asp Arg Val
 1          5          10          15
Glu Asp Thr Leu Ala Ala His Arg Asn Glu Leu Val Ser Leu Leu Ser
          20          25          30
Arg Tyr Val Ala Gln Gly Lys Gly Ile Leu Gln Pro His His Leu Leu
          35          40          45
Asp Glu Leu Glu Asn Ile Ile Ser Glu Asp Glu Gly Lys Ser Ser Leu
          50          55          60
Ser Asp Gly Pro Phe Ser
65          70

```

<210> 103  
 <211> 534  
 <212> PRT  
 <213> Eucalyptus grandis

<400> 103  
 Val Leu Phe Thr His Leu Pro Pro Gln Lys Pro Asn Arg Ile Ser Leu  
 1 5 10 15  
 Leu Leu Phe Phe Ile Phe His Ile Thr Thr Phe Leu Leu Leu Leu Leu  
 20 25 30  
 Leu Leu Ser Val Leu Ser Thr Phe Ile Ser Ile Ala Val Ser Leu Ser  
 35 40 45  
 Asp Pro Glu Leu Phe Phe Ala Ser Pro Pro Met Ala Ala Ala Thr  
 50 55 60  
 Leu Ser Ala Pro Asp Ala Ala Lys Leu Ser Gln Leu Lys Ser Ala Val  
 65 70 75 80  
 Ser Gly Leu Gly Gln Ile Ser Glu Ser Glu Lys Asn Gly Phe Ile Asn  
 85 90 95  
 Leu Val Ser Arg Tyr Leu Ser Gly Glu Ala Gln His Val Asp Trp Ser  
 100 105 110  
 Lys Ile Gln Thr Pro Thr Asp Glu Ile Val Val Pro Tyr Asp Ser Leu  
 115 120 125  
 Ala Pro Thr Pro Gln Asp Pro Ala Ala Thr Lys Ser Leu Leu Asp Lys  
 130 135 140  
 Leu Val Val Leu Lys Leu Asn Gly Gly Leu Gly Thr Thr Met Gly Cys  
 145 150 155 160  
 Thr Gly Pro Lys Ser Val Ile Glu Val Arg Asn Gly Leu Thr Phe Leu  
 165 170 175  
 Asp Leu Ile Val Ile Gln Ile Glu Asn Leu Asn Thr Lys Tyr Gly Cys  
 180 185 190  
 Asn Val Pro Leu Leu Leu Met Asn Ser Phe Asn Thr His Asp Asp Thr  
 195 200 205  
 Leu Lys Ile Val Glu Lys Tyr Ala Asn Ser Asn Ile Asp Ile His Thr  
 210 215 220  
 Phe Asn Gln Ser Gln Tyr Pro Arg Leu Val Val Glu Asp Phe Met Pro  
 225 230 235 240  
 Leu Pro Cys Lys Gly Gln Thr Gly Lys Asp Gly Trp Tyr Pro Pro Gly  
 245 250 255  
 His Gly Asp Val Phe Ala Ser Leu Met Asn Ser Gly Lys Leu Asp Ala  
 260 265 270  
 Leu Leu Ser Gln Gly Lys Glu Tyr Val Phe Ala Ala Asn Ser Asp Asn  
 275 280 285  
 Leu Gly Ala Ile Val Asp Leu Lys Ile Leu Asn His Leu Met Thr Asn  
 290 295 300  
 Lys Asn Glu Tyr Cys Met Glu Val Thr Pro Lys Thr Leu Ala Asp Val  
 305 310 315 320  
 Lys Gly Gly Thr Leu Ile Ser Tyr Glu Gly Lys Val Gln Leu Leu Glu  
 325 330 335  
 Ile Ala Gln Val Pro Asp Glu His Ile Asn Glu Phe Lys Ser Ile Glu  
 340 345 350  
 Lys Phe Lys Ile Phe Asn Thr Asn Asn Leu Trp Val Asn Leu Lys Ala  
 355 360 365  
 Ile Lys Arg Leu Val Glu Ala Gln Ala Leu Lys Met Glu Ile Ile Pro  
 370 375 380  
 Asn Pro Lys Glu Val Asp Gly Ile Lys Val Leu Gln Leu Glu Thr Ala  
 385 390 395 400  
 Ala Gly Ala Ala Ile Lys Phe Phe Asp Asn Ala Ile Gly Ile Asn Val

405 410 415  
 Pro Arg Ser Arg Phe Leu Pro Val Lys Ala Thr Ser Asp Leu Leu Leu  
 420 425 430  
 Val Gln Ser Asp Leu Tyr Thr Leu Val Asp Gly Phe Val Glu Arg Asn  
 435 440 445  
 Lys Ala Arg Thr Asn Pro Ser Asn Pro Ser Ile Glu Leu Gly Pro Glu  
 450 455 460  
 Phe Lys Lys Val Gly Asn Phe Leu Ser Arg Phe Lys Ser Ile Pro Ser  
 465 470 475 480  
 Ile Ile Glu Leu Asp Ser Leu Lys Val Ser Gly Asp Val Trp Phe Gly  
 485 490 495  
 Thr Gly Ile Thr Leu Lys Gly Lys Val Thr Ile Ala Ala Lys Pro Gly  
 500 505 510  
 Val Lys Leu Glu Ile Pro Asp Gly Val Val Leu Glu Asn Lys Glu Ile  
 515 520 525  
 His Gly Pro Glu Asp Leu  
 530

<210> 104  
 <211> 480  
 <212> PRT  
 <213> Pinus radiata

<400> 104  
 Met Ala Ala Ala Pro Ala Val Ala Ser Pro Ala Ala Glu Thr Asp Arg  
 1 5 10 15  
 Ile Pro Lys Leu Gln Ala Glu Val Thr Lys Leu Asn Gln Ile Ser Asp  
 20 25 30  
 Asn Glu Lys Glu Gly Phe Val Arg Leu Val Ser Arg Tyr Leu Ser Gly  
 35 40 45  
 Glu Glu Glu Lys Ile Glu Trp Glu Lys Ile Lys Thr Pro Thr Asp Glu  
 50 55 60  
 Ile Val Val Pro Tyr Asp Thr Leu Ala Ala Leu Gly Glu Asp Pro Ser  
 65 70 75 80  
 Glu Thr Lys Glu Leu Leu Asp Lys Leu Val Val Leu Lys Leu Asn Gly  
 85 90 95  
 Gly Leu Gly Thr Thr Met Gly Cys Thr Gly Pro Lys Ser Val Ile Glu  
 100 105 110  
 Val Arg Asn Gly Leu Thr Phe Leu Asp Leu Ile Val Lys Gln Ile Glu  
 115 120 125  
 Ser Leu Asn Asn Lys Tyr Asp Ser Lys Val Pro Leu Val Leu Met Asn  
 130 135 140  
 Ser Phe Asn Thr His Asp Asp Thr Ile Lys Ile Val Glu Lys Tyr Ser  
 145 150 155 160  
 Gly Ser Asn Ile Asp Ile His Ile Phe Asn Gln Ser Gln Tyr Pro Arg  
 165 170 175  
 Met Val Ala Glu Asp Leu Thr Pro Trp Pro Thr Lys Gly Arg Thr Asp  
 180 185 190  
 Lys Glu Ala Trp Tyr Pro Pro Gly His Gly Asp Val Phe Pro Ala Leu  
 195 200 205  
 Leu Asn Ser Gly Lys Leu Asp Glu Leu Leu Ser Gln Gly Lys Glu Tyr  
 210 215 220  
 Val Phe Ile Ala Asn Ser Asp Asn Leu Gly Ala Ile Val Asp Leu Lys  
 225 230 235 240  
 Ile Leu Asn His Leu Val Lys Asn Lys Asn Glu Tyr Cys Met Glu Val  
 245 250 255  
 Thr Pro Lys Thr Leu Ala Asp Val Lys Gly Gly Thr Leu Ile Ser Tyr

260 265 270  
 Glu Gly Arg Val Gln Leu Leu Glu Ile Ala Gln Val Pro Glu Glu His  
 275 280 285  
 Val Gly Glu Phe Lys Ser Ile Glu Lys Phe Lys Ile Phe Asn Thr Asn  
 290 295 300  
 Asn Leu Trp Val Asn Leu Lys Ala Ile Lys Arg Leu Val Glu Ala Asp  
 305 310 315 320  
 Ala Leu Lys Met Glu Ile Ile Pro Asn Pro Lys Glu Val Asp Gly Val  
 325 330 335  
 Lys Val Leu Gln Leu Glu Thr Ala Ala Gly Ala Ala Ile Arg Phe Phe  
 340 345 350  
 Asp Arg Ala Ile Gly Val Asn Val Pro Arg Ser Arg Phe Leu Pro Val  
 355 360 365  
 Lys Ala Thr Ser Asp Leu Leu Leu Val Gln Ser Asp Leu Tyr Thr Val  
 370 375 380  
 Glu Glu Gly Phe Val Ile Arg Asn Pro Ala Arg Val Asn Pro Thr Asn  
 385 390 395 400  
 Pro Thr Ile Glu Leu Gly Pro Glu Phe Lys Lys Val Gly Asn Phe Leu  
 405 410 415  
 Lys Arg Phe Lys Ser Ile Pro Ser Ile Ile Asp Leu Asp Ser Leu Lys  
 420 425 430  
 Val Ser Gly Asp Val Trp Phe Gly Ser Gly Val Ile Leu Lys Gly Lys  
 435 440 445  
 Val Ile Ile Glu Ala Lys Gln Gly Ala Thr Leu Glu Ile Pro Asp Glu  
 450 455 460  
 Ser Val Ile Glu Asn Lys Val Val Ser Ser Pro Asp Asp Ile Val Asn  
 465 470 475 480

&lt;210&gt; 105

&lt;211&gt; 573

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 105

ctcactcgat	ctcgaaggcc	agaaggggga	ggccgagcct	cttgcttttt	ttcgtgtata	60
aaagggcctc	ccccattcct	catttttcac	catcctccgt	tcgttcgttc	ccttcccttt	120
ccattgttgc	gtttaagccc	tccaattttc	ttttggcgtc	ccgttttttg	ggctcccttg	180
aagatctcct	cttcatttcg	ggatttcctg	ccttcgccgc	gccatttgaa	gttctttttc	240
tgagagaaga	atthagacat	ggctgatcgc	atgctgactc	gaagccacag	ccttcgcgag	300
cgtttgagac	agaccctctc	tgctcaccgc	aacgatattg	tggccttcct	ttcaagggtt	360
gaagccaagg	gcaaaggcat	cttgacgcgc	caccagatct	ttgctgagtt	tgaggccatc	420
tctgaggaga	gcagagcaaa	gcttcttgat	ggggcctttg	gtgaagtcct	caaatccact	480
caggaagcga	ttgtgtcgcc	tccatgggtt	gctcttgctg	ttcgtccaag	gccggggcgtg	540
tgggagcaca	tccgtgtgaa	cgtccatgcg	ctt			573

&lt;210&gt; 106

&lt;211&gt; 105

&lt;212&gt; PRT

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 106

Met Ala Asp Arg Met Leu Thr Arg Ser His Ser Leu Arg Glu Arg Leu

```

      1           5           10           15
Asp Glu Thr Leu Ser Ala His Arg Asn Asp Ile Val Ala Phe Leu Ser
      20           25           30
Arg Val Glu Ala Lys Gly Lys Gly Ile Leu Gln Arg His Gln Ile Phe
      35           40           45
Ala Glu Phe Glu Ala Ile Ser Glu Glu Ser Arg Ala Lys Leu Leu Asp
      50           55           60
Gly Ala Phe Gly Glu Val Leu Lys Ser Thr Gln Glu Ala Ile Val Ser
      65           70           75           80
Pro Pro Trp Val Ala Leu Ala Val Arg Pro Arg Pro Gly Val Trp Glu
      85           90           95
His Ile Arg Val Asn Val His Ala Leu
      100           105

```

<210> 107  
 <211> 664  
 <212> DNA  
 <213> Eucalyptus grandis

```

<400> 107
ggcagcagct cttctcgtct cgctttctca tataaagaag tgaaagaata cgaggatact      60
ccacttgggt atcgccaaga actcattggg tcgcgagaag attggccaac atgatggaat      120
ccgggggttc cctgtgcaac acttgcgagg aggcgtgttg ggttgatgag aaaggcgagg      180
tcttcgtggc ttgtcaagag tgcaacttcg ccatttgcaa ggcttggtgc gaatatgaga      240
ttaaggaagg aagaaaagcg tgcttgcgct gtggcactcc atttgaagcg aactcgatgg      300
ctgatgctga gagaaatgaa ttgggaagtc gatcgacaat ggcagctcaa ctcaatgatc      360
ctcaggacac agggattcat gctagacaca tcagcagtgt ttctacgttg gatagtgaat      420
acaatgatga gactgggaac cctatctgga agaataagag ggagagctgg aaggacaaaa      480
agaataagaa gaagaaggcc ccgacgaagg ctgagaaaaga ggctcaagtt ccaccagagc      540
agcagatgga agagaagcaa attgctgatg cttcagagcc actctcgacc gttattccca      600
ttgccaaaag caaactcgca ccataccgaa ccgtaataat aatgcgattg atcattttgg      660
cact

```

<210> 108  
 <211> 184  
 <212> PRT  
 <213> Eucalyptus grandis

```

<400> 108
Met Met Glu Ser Gly Val Pro Leu Cys Asn Thr Cys Gly Glu Ala Val
      1           5           10           15
Gly Val Asp Glu Lys Gly Glu Val Phe Val Ala Cys Gln Glu Cys Asn
      20           25           30
Phe Ala Ile Cys Lys Ala Cys Val Glu Tyr Glu Ile Lys Glu Gly Arg
      35           40           45
Lys Ala Cys Leu Arg Cys Gly Thr Pro Phe Glu Ala Asn Ser Met Ala
      50           55           60
Asp Ala Glu Arg Asn Glu Leu Gly Ser Arg Ser Thr Met Ala Ala Gln
      65           70           75           80
Leu Asn Asp Pro Gln Asp Thr Gly Ile His Ala Arg His Ile Ser Ser
      85           90           95
Val Ser Thr Leu Asp Ser Glu Tyr Asn Asp Glu Thr Gly Asn Pro Ile
      100           105           110
Trp Lys Asn Arg Val Glu Ser Trp Lys Asp Lys Lys Asn Lys Lys Lys
      115           120           125
Lys Ala Pro Thr Lys Ala Glu Lys Glu Ala Gln Val Pro Pro Glu Gln
      130           135           140

```

Gln Met Glu Glu Lys Gln Ile Ala Asp Ala Ser Glu Pro Leu Ser Thr  
 145 150 155 160  
 Val Ile Pro Ile Ala Lys Ser Lys Leu Ala Pro Tyr Arg Thr Val Ile  
 165 170 175  
 Ile Met Arg Leu Ile Ile Leu Ala  
 180

&lt;210&gt; 109

&lt;211&gt; 1293

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 109

ctgactctct	ctctctctgt	tttgtctcct	ccctcctctc	tctcgttttc	gcttcgtcgt	60
gaacgcaccc	acacgatctt	ccattccctc	aacaatgtcg	actctcaccc	tcccgcagcc	120
actgccccct	gtagccgatg	actgcgagca	gctccggaca	gccttcgcag	gatggggaac	180
aaatgagaag	ctgatcatat	ccatattggg	tcataggaat	gcggcgcaga	ggaagctgat	240
tcggcaaaacc	tatgccgaga	cttacggcga	ggacctcctc	aaggcattgg	acagagaact	300
taccaatgat	ttcgagaggc	tggtggctct	ttggtcactt	gatccggctg	aacgtgatgc	360
gtacttggcg	aatgaagcga	cgaaaagatg	gacttcaagc	aaccaggttc	tcatggaaat	420
agcctgcacg	aggctctccg	agcagttgct	tatggcaaga	caagcatatc	atgcccgata	480
caagaagtca	atggaagagg	acgtcgtctc	ccacacaact	ggagattttc	gtaagttgct	540
ggtacctctt	gggagctcct	accgtaatga	tggagatgag	gtgaatatga	ctttggcaaa	600
agcagaggct	aagatactcc	acgagaagat	ctcagagaag	gcttatggcc	atgaggatct	660
cataaggatt	ttggctacta	ggagcaaaagc	acaggtcaat	gctacgctga	atcactacaa	720
aaatgagttt	ggaaatgata	tcaacaagga	tttgaaaact	gatccaaaag	acgcgttctt	780
tactatactg	agagctacag	taaagtgcct	gactcgcctt	gagaagtatt	ttgaaaaggt	840
tcttcgtcta	gccatcaata	agcgagggaac	agatggaagg	gctctgacca	gagtagttgc	900
taccagggcc	gaggttgaca	tgaagtttat	aagtgaggag	taccagagga	ggaatagcat	960
ccctctcgat	cgtgccattg	tcaaggacac	tactggagac	tatgaaaaaa	tgcttctggc	1020
attgattggc	cacgtcgagg	cttgatttac	aagtactcat	gaagctatcc	tggtggaggc	1080
aatatctctg	tttttgggtg	ggtttgaggc	atttctatct	tccttgcttt	ccaacaacgt	1140
gtagttacca	acatgcctcc	ccagttgtca	gttgtagcta	tgcgaagcaa	atacacttct	1200
tataatggcg	ttgggttatg	tacttatgag	aagtctttga	ttttgatctt	taatcaagac	1260
tgctagtaag	tgatcgtgaa	aaaaaaaaaa	aaa			1293

&lt;210&gt; 110

&lt;211&gt; 484

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 110

ggaagctgat	tcggcaaaacc	tatgccgaga	cttacggcga	ggacctcctc	aaggcattgg	60
acagagaact	taccaatgat	tttgaggtct	gatcttcttt	aattatttgt	attcatccca	120
tggagacgcg	tccctctttc	tctcagatta	atccatattc	attccgtatc	gtcagaggct	180
ggtggctcct	tggtcgcttg	atccggctga	acgtgatgcg	tacttggcga	atgaagcgac	240
gaaaagatgg	acttcaagca	accaggttct	catggaaata	gcctgcacga	ggtctccaca	300
gcagttgctc	atggcaagac	aagcatatca	tgctcgatac	aagaagtcgc	tggaagagga	360
cgctcgtcac	cacacaactg	gagattttcg	taagttgctg	gtacctcttg	tgagctccta	420
ccattatgat	ggagatgagg	tgaatatgac	tttggcaaaa	gcagaggcta	agatactcca	480
cgag						484

&lt;210&gt; 111

&lt;211&gt; 221

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 111

cgtacttggc	gaatgaagcg	acgaaaagat	ggacttcaag	caaccagggtt	ctaattggaaa	60
tagcctgcac	gaggtctccg	cagcagttgc	ttatggcaag	acaagcatat	catgccccgat	120
acaagaagtc	gctggaagag	gacgtcggtc	accacacaac	tggagatttt	cgtaagtgtgc	180
tggtacctct	tgtgagctcc	taccgttatg	atggagatga	g		221

&lt;210&gt; 112

&lt;211&gt; 789

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 112

atcgtcttcg	gctcctcgcg	atatcaccaa	cttgcttccg	cacagagaga	gagagagaga	60
gagagagaga	gaatggcgac	tatcgcggtg	ccaccgtcgg	ttccgtctcc	ggctgaggat	120
gccgagcagc	tccaaaaagc	tttcgcagga	tgggggacga	atgaagatct	gatcatatcc	180
atactgcctc	acagaaacgc	agcgcagcgg	aaagtaatcc	gacaaacata	tgccgagaca	240
tatggggaag	atcttctcaa	agcgttgac	aaggaactct	ctagtgactt	tgagagatct	300
gtgcttctgt	ggaccctgga	tcctgcggag	cgtgatgcat	tcttgtccaa	tgaagctacc	360
aagagattga	cttcgagcaa	ctgggttctc	atggaaattg	cttgcacgag	gtcttcaatg	420
gagttattca	tggtgaggca	ggcctatcat	gctcgttata	agaaatctct	tgaagaagac	480
atcgcatatc	acactactgg	ggatttccgc	aagctgcttg	ttcctctggc	aagtaccttt	540
cggtatgagg	ggcctgaggt	gaacatgaca	ttggcgagat	cagaggctaa	gatacttcat	600
gagaagattc	acgagaaggc	ttacaatcat	gatgagctca	tcagaattgt	tactacaaga	660
agtaaagctc	agcttaatgc	aaccctcaat	tactacaaca	atgagtttgg	gaatgccatc	720
aacaaggatc	tgaaggctga	tccaaatgat	gaatttctga	aactgctgag	atcagcaatt	780
aagtgccttg						789

&lt;210&gt; 113

&lt;211&gt; 704

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 113

gttttgttga	gctactagat	tttagtaaat	caagaattca	tcagctataa	attgaggcat	60
tcgatttcag	ttttagttac	atthttgtga	agttgggtcga	cctgcattgc	tgaagatattc	120
gtgcgaagta	tgtgatttgt	cgagaagatg	tcaacaatta	tagtgccagt	tccaataaccg	180
accccatctg	aagactctga	acgcctgagg	aaggcttttg	aagggtgggg	cacaaatgag	240
aagtcaatca	tacaaatatt	aggacataga	actgcagcac	aacgcaaagt	aatccgtcaa	300
agttattttc	aactgtacga	agaggatctc	ttgaagcgat	tagaatctga	gctttcaagt	360
gactttgaga	aagctgtatt	cctttgggta	ctagatccag	ctgaacgtga	tgcggtcata	420
tctcatgggtg	caataaagaa	gtggaatgca	agaatataat	cgctttttaga	aattttccagt	480
gctcgatctt	cggctgaact	attgatgggtg	aggcaagcat	atcatattcg	gtacaaaaag	540
tccctcgaag	aagacgtggc	tgacataaca	agtggaaaact	tccgtaagtt	gctggttagca	600
cttgtaagtt	catatcggtg	tgaagggtccg	gaagtggata	tgcatattggc	aagttatgaa	660
gcaaagaagc	taagtgaatc	tataaccgag	caaaaaagat	aatt		704

&lt;210&gt; 114

&lt;211&gt; 316

&lt;212&gt; PRT

&lt;213&gt; Pinus radiata

&lt;400&gt; 114

Met	Ser	Thr	Leu	Thr	Val	Pro	Gln	Pro	Leu	Pro	Pro	Val	Ala	Asp	Asp
1			5					10					15		
Cys	Glu	Gln	Leu	Arg	Thr	Ala	Phe	Ala	Gly	Trp	Gly	Thr	Asn	Glu	Lys
			20					25					30		
Leu	Ile	Ile	Ser	Ile	Leu	Gly	His	Arg	Asn	Ala	Ala	Gln	Arg	Lys	Leu



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      35      40      45
Ile Arg Gln Thr Tyr Ala Glu Thr Tyr Gly Glu Asp Leu Leu Lys Ala
 50      55      60
Leu Asp Arg Glu Leu Thr Asn Asp Phe Glu Arg Leu Val Val Leu Trp
 65      70      75      80
Ser Leu Asp Pro Ala Glu Arg Asp Ala Tyr Leu Ala Asn Glu Ala Thr
      85      90      95
Lys Arg Trp Thr Ser Ser Asn Gln Val Leu Met Glu Ile Ala Cys Thr
      100      105      110
Arg Ser Pro Gln Gln Leu Leu Met Ala Arg Gln Ala Tyr His Ala Arg
      115      120      125
Tyr Lys Lys Ser Met Glu Glu Asp Val Ala His His Thr Thr Gly Asp
      130      135      140
Phe Arg Lys Leu Leu Val Pro Leu Gly Ser Ser Tyr Arg Asn Asp Gly
 145      150      155      160
Asp Glu Val Asn Met Thr Leu Ala Lys Ala Glu Ala Lys Ile Leu His
      165      170      175
Glu Lys Ile Ser Glu Lys Ala Tyr Gly His Glu Asp Leu Ile Arg Ile
      180      185      190
Leu Ala Thr Arg Ser Lys Ala Gln Val Asn Ala Thr Leu Asn His Tyr
      195      200      205
Lys Asn Glu Phe Gly Asn Asp Ile Asn Lys Asp Leu Lys Thr Asp Pro
      210      215      220
Lys Asp Ala Phe Leu Thr Ile Leu Arg Ala Thr Val Lys Cys Leu Thr
 225      230      235      240
Arg Pro Glu Lys Tyr Phe Glu Lys Val Leu Arg Leu Ala Ile Asn Lys
      245      250      255
Arg Gly Thr Asp Glu Gly Ala Leu Thr Arg Val Val Ala Thr Arg Ala
      260      265      270
Glu Val Asp Met Lys Phe Ile Ser Glu Glu Tyr Gln Arg Arg Asn Ser
      275      280      285
Ile Pro Leu Asp Arg Ala Ile Val Lys Asp Thr Thr Gly Asp Tyr Glu
      290      295      300
Lys Met Leu Leu Ala Leu Ile Gly His Val Glu Ala
 305      310      315

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<210> 115  
 <211> 111  
 <212> PRT  
 <213> Pinus radiata

```

      <400> 115
Ser Ile Phe Ile Pro Tyr Arg Gln Arg Leu Val Val Leu Trp Ser Leu
 1      5      10      15
Asp Pro Ala Glu Arg Asp Ala Tyr Leu Ala Asn Glu Ala Thr Lys Arg
      20      25      30
Trp Thr Ser Ser Asn Gln Val Leu Met Glu Ile Ala Cys Thr Arg Ser
      35      40      45
Pro Gln Gln Leu Leu Met Ala Arg Gln Ala Tyr His Ala Arg Tyr Lys
      50      55      60
Lys Ser Leu Glu Glu Asp Val Ala His His Thr Thr Gly Asp Phe Arg
 65      70      75      80
Lys Leu Leu Val Pro Leu Val Ser Ser Tyr His Tyr Asp Gly Asp Glu
      85      90      95
Val Asn Met Thr Leu Ala Lys Ala Glu Ala Lys Ile Leu His Glu
      100      105      110

```

<210> 116  
 <211> 73  
 <212> PRT  
 <213> Pinus radiata

<400> 116  
 Tyr Leu Ala Asn Glu Ala Thr Lys Arg Trp Thr Ser Ser Asn Gln Val  
 1 5 10 15  
 Leu Met Glu Ile Ala Cys Thr Arg Ser Pro Gln Gln Leu Leu Met Ala  
 20 25 30  
 Arg Gln Ala Tyr His Ala Arg Tyr Lys Lys Ser Leu Glu Asp Val  
 35 40 45  
 Gly His His Thr Thr Gly Asp Phe Arg Lys Leu Leu Val Pro Leu Val  
 50 55 60  
 Ser Ser Tyr Arg Tyr Asp Gly Asp Glu  
 65 70

<210> 117  
 <211> 239  
 <212> PRT  
 <213> Pinus radiata

<400> 117  
 Met Ala Thr Ile Ala Val Pro Pro Ser Val Pro Ser Pro Ala Glu Asp  
 1 5 10 15  
 Ala Glu Gln Leu Gln Lys Ala Phe Ala Gly Trp Gly Thr Asn Glu Asp  
 20 25 30  
 Leu Ile Ile Ser Ile Leu Pro His Arg Asn Ala Ala Gln Arg Lys Val  
 35 40 45  
 Ile Arg Gln Thr Tyr Ala Glu Thr Tyr Gly Glu Asp Leu Leu Lys Ala  
 50 55 60  
 Leu Asp Lys Glu Leu Ser Ser Asp Phe Glu Arg Ser Val Leu Leu Trp  
 65 70 75 80  
 Thr Leu Asp Pro Ala Glu Arg Asp Ala Phe Leu Ser Asn Glu Ala Thr  
 85 90 95  
 Lys Arg Leu Thr Ser Ser Asn Trp Val Leu Met Glu Ile Ala Cys Thr  
 100 105 110  
 Arg Ser Ser Met Glu Leu Phe Met Val Arg Gln Ala Tyr His Ala Arg  
 115 120 125  
 Tyr Lys Lys Ser Leu Glu Glu Asp Ile Ala Tyr His Thr Thr Gly Asp  
 130 135 140  
 Phe Arg Lys Leu Leu Val Pro Leu Ala Ser Thr Phe Arg Tyr Glu Gly  
 145 150 155 160  
 Pro Glu Val Asn Met Thr Leu Ala Arg Ser Glu Ala Lys Ile Leu His  
 165 170 175  
 Glu Lys Ile His Glu Lys Ala Tyr Asn His Asp Glu Leu Ile Arg Ile  
 180 185 190  
 Val Thr Thr Arg Ser Lys Ala Gln Leu Asn Ala Thr Leu Asn Tyr Tyr  
 195 200 205  
 Asn Asn Glu Phe Gly Asn Ala Ile Asn Lys Asp Leu Lys Ala Asp Pro  
 210 215 220  
 Asn Asp Glu Phe Leu Lys Leu Leu Arg Ser Ala Ile Lys Cys Leu  
 225 230 235

<210> 118  
 <211> 184  
 <212> PRT

<213> *Pinus radiata*

&lt;400&gt; 118

```

Met Ser Thr Ile Ile Val Pro Val Pro Ile Pro Thr Pro Ser Glu Asp
 1           5           10           15
Ser Glu Arg Leu Arg Lys Ala Phe Glu Gly Trp Gly Thr Asn Glu Lys
      20           25           30
Ser Ile Ile Gln Ile Leu Gly His Arg Thr Ala Ala Gln Arg Lys Val
      35           40           45
Ile Arg Gln Ser Tyr Phe Gln Leu Tyr Glu Glu Asp Leu Leu Lys Arg
 50           55           60
Leu Glu Ser Glu Leu Ser Ser Asp Phe Glu Lys Ala Val Phe Leu Trp
65           70           75           80
Val Leu Asp Pro Ala Glu Arg Asp Ala Val Ile Ser His Gly Ala Ile
      85           90           95
Lys Lys Trp Asn Ala Lys Asn Ile Ser Leu Leu Glu Ile Ser Ser Ala
      100          105          110
Arg Ser Ser Ala Glu Leu Leu Met Val Arg Gln Ala Tyr His Ile Arg
      115          120          125
Tyr Lys Lys Ser Leu Glu Glu Asp Val Ala Ala His Thr Ser Gly Asn
130          135          140
Phe Arg Lys Leu Leu Val Ala Leu Val Ser Ser Tyr Arg Tyr Glu Gly
145          150          155          160
Pro Glu Val Asp Met His Leu Ala Ser Tyr Glu Ala Lys Lys Leu Ser
      165          170          175
Glu Ser Ile Thr Glu Gln Lys Arg
      180

```

&lt;210&gt; 119

&lt;211&gt; 568

&lt;212&gt; DNA

<213> *Eucalyptus grandis*

&lt;400&gt; 119

```

tcgtcacc ca attcctcacc aacaacaacc gagcctctgg cacactttcc tccatcagga      60
ggttctacgt ccaggacggc aaagtaattc caaactctat ggtaaaccctc tccgggtcttc      120
ccaaagtcaa ctcgatcacg tcagattact gcaccgctaa aatggacggt ctcgacgatt      180
ctaccgcttt caacgtacat ggtggtcttg caaagatggg taaatccctt gcacgaggag      240
cagtactcgt ggtcagtcctc tgggatgatc ttggcgggcg gatgacttgg ttggatgggc      300
tagcagggga tgcattctgcc cctgggaccc tccgtggacc gtgcaccgct gcgaatgtaa      360
catcagatcc ggctacctcc gtcactttct cgaatatccg agttggcgat atcaatagca      420
ctttctctca ggtgcacttt gggcaatgtg gaggtcaatt ttacgagggg ccagctcttt      480
gcgcagaccc attcgagtgt gtcttcagta atccgtatta cagccagtgt ctataagata      540
ttgaatataa cagggtttat gtccttcc

```

&lt;210&gt; 120

&lt;211&gt; 360

&lt;212&gt; DNA

<213> *Eucalyptus grandis*

&lt;400&gt; 120

cgacatgttc	gacaaggccg	ctctcttcgc	tttctcttta	ctcgccgtca	cttacgggtca	60
gcaggtcggt	acccagactg	ctgaatctca	tccgcctctt	acctggcaaa	aatgtacgac	120
tgctgggtgga	tgtaccaatg	tttctgggtg	tagtggtgtc	attgacgcga	attggcggtg	180
ggtccattcc	atcaacggta	ctactaactg	ttacactggg	caagcatgga	acacgacact	240
ctgtccggat	gacacgactt	gcgcagccaa	ctgcgctttg	gatgggtgctg	attactctgg	300
cacttatggc	attactactt	ccggcaatgc	tcttaccctc	aagttcgtca	ctcaatcttc	360

&lt;210&gt; 121

&lt;211&gt; 375

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 121

tattatgatg	cgaatttaaat	tgctatagca	gttgggttta	gcaggacaat	ttacagtgtg	60
attcctcaat	ggagccgttt	gataggtggg	gtcttcttca	gtttttgggt	cttggtcat	120
ctgtatcctt	ttgcaaagg	gctcatggga	agacgtgggc	gcacccctac	cattgttttc	180
gtttgggtcag	gactcattgc	aatcaccata	tcacttcttt	gggtggcaat	cagcccccca	240
gctgggtcaa	cccaaatcgg	tggtcttttc	cagttccctt	gataggttat	tctttttaat	300
atgctttatc	tgtttagtga	cattactcca	ttctttttta	aatgagatga	tcattgtgac	360
aaaaaaaaaa	aaaaa					375

&lt;210&gt; 122

&lt;211&gt; 590

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 122

cacgactttg	gaagatgggtg	gtgttccgcc	agatgctagt	cctgcatcgc	tactaaaaga	60
agccatccaa	gtcatcagtt	gcggatatga	agacaagaca	gaatggggaa	aagaagtggg	120
ctggatatac	ggttcgggtga	ctgaggatat	attgactggg	ttcaaaatgc	actgccacgg	180
ctggagatcg	gtgtactgta	tacctaaagag	gcctgcattc	aagggttcag	caccgatcaa	240
cctttcggat	cgtctacacc	aggttctccg	gtgggctctt	gggtcagttg	agattttctt	300
gagcaggcat	tgcccaatct	ggtatggcta	cggggagggt	ttgaagtggg	tggaaacgatt	360
ttcttacatc	aattcgggtg	tgtatccttg	gacctccatc	cccttgattg	tttactgctc	420
actccgggct	atctgccttc	tcactggcca	attcatcgtg	cctgagatta	gcaactatgc	480
aagtctcgtc	tttatggcac	ttttcatctc	cattgctgcg	actggtatcc	ttgagatgca	540
atgggggtgg	gttggaatcg	atgactgggtg	gagaaacgag	cagttttggg		590

&lt;210&gt; 123

&lt;211&gt; 590

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 123

cactgcaactg	attcctaaga	agttcggaaa	ctcatatcatg	ttcattgatt	ccataccttt	60
agctgagttc	caaggccgac	cccttgccga	ccatccatcc	gtgaaaaatg	gacggcctcc	120
tggtgctctc	actgttcttc	gacggcttct	tgatgcgtca	acagtgtcag	aggcaataag	180
tgccatctca	tgctggtagc	aagataaaac	tgagtggggg	gaacgagtag	gggtggattta	240
cgggtccgctc	acagaggatg	tcgtgacagg	gtatagaatg	cacaatcgag	gatggacatc	300
tgtatactgc	gtgaccaaga	gagatgcttt	tcgtggggacc	gcacccatca	atctcaccca	360
tcgggttcat	caagtccctg	gctggggcag	gggctcggtg	gagatattct	tctctcgcaa	420
caatgccata	atggccagcg	gcaggctgaa	gttccttcag	aggattgctt	acctcaacgt	480
tggaaatttac	cctttcactt	ccatctttct	tattgtctac	tgctttctcc	cggcgctctc	540
tctattctct	gggaagtcca	ttgtgcaatc	gctcagtgta	tccttcctaa		590

&lt;210&gt; 124

&lt;211&gt; 619

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 124

ggaaaggggtg	gtgacaagaa	ttacatcgac	aagaagagag	ctggcaaaaag	aactgaatcc	60
aacatttccaa	tattcaacat	ggaggatatt	gaggagggga	tggaagggtta	tgatgatgag	120
agatcactgc	ttatgtccca	gaaaagctta	gagaagcgct	ttgggtcaatc	gccagtcttc	180
attgcagcaa	cattcatgga	acagggaggg	cttccacat	ctactaatcc	agcaactctt	240
ttgaaggaag	caattcatgt	tatcagctgt	ggctatgagg	acaagactga	gtggggcaaa	300
gagattggat	ggatatacgg	ttctgtcaca	gaagatatct	taacaggggt	taagatgcat	360
gctcgagggt	ggatctccat	ttactgtatg	cctccacgcc	cagcattcaa	gggttccgct	420
cccataaatc	tttcagatcg	tctgaaccaa	gttcttcgat	gggcattggg	gtccattgag	480
atthttactaa	gcaggcattg	tcctatatgg	tatgggttaca	atgggagact	gaagtgggtg	540
gagagattgg	catatataaa	taccattgtg	tatccctca	cttcaatcct	cttgattgct	600
tattgcatc	tgcttgc					619

&lt;210&gt; 125

&lt;211&gt; 429

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 125

cgctctctta	gccccttgcc	cagcagttgc	aagtcacatca	ttcatcatca	tgctcctcca	60
gactggcgct	ttcctcgcca	cgctcctggg	caccgccc	gcccaggccg	taggcaagga	120
gcagaccgag	actcacccca	agatgacatg	gaagaagtgc	agtagcggtg	gcagctgcac	180
tagcgtgaac	ggtgaagtca	ccatcgacgc	caattggcgc	tggtctgcacg	ggacgtcaga	240
caccaagaac	tgctacgatg	gcaacaagtg	gaccgacaag	tgacgacg	cgactgactg	300
cgccctccaag	tgccgcatcg	aaggtgccac	ctattccaag	acatacgggtg	cctccactag	360
cggtgatgcc	ctgactctca	agttcgctcac	caagcacgaa	tatgggtacca	atatacggctc	420
tcgtctcta						429

&lt;210&gt; 126

&lt;211&gt; 534

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 126

ctggaatcat	cctttatttc	tgtgactgtg	cactttcaga	gaggaaggag	agagaggaaa	60
cccaaaagaa	aaggttgtgg	cttgcagctg	aatcaatacc	acacacccat	atatacaata	120
ctcccacact	attccctttt	tttcttctt	cattaatttt	atctctcttc	atthttgtaat	180
ttagatatth	ccaaacaaaa	gtctgtctct	ttttttcttt	tattattatc	atthttgtccg	240
actccgattt	gccgtttgag	agaagttacc	tctgttatgg	actgtggatc	tccgagaacc	300
aagctgtcct	gagttgaccc	tggtgtctct	tttaacccac	tcaaccgatc	tgtcaagatt	360
gtagcctctg	tggtccgaca	aatgaatacc	gggtgtcggt	taatcgccgg	ttcacacaa	420
aggaacgagt	ttgttcttat	taatgccgat	gagaatgcc	gaataagatc	agtgcaagag	480
ctgagtgggc	agacgtgtca	aatctgtaga	gacgagattg	aattgaccgt	cgat	534

&lt;210&gt; 127

&lt;211&gt; 450

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 127

aaagcactga	gtgagagctg	gaactgaagt	gactgactga	tgtagagag	agagagaatt	60
gagatagaga	tgagtgacg	aggaagcctc	ccctcccttc	ttcaccaaac	gttcgctctc	120
tcctcgctcca	cactccttc	gctgctgccc	cctccattgc	gtagcaccgt	cgccgcccgt	180
cgccgcccgt	ctcctcttct	ccgagacccg	gaatcgcgaa	ccgcttgctg	agcaccgcga	240

```

tcgccccga gcgagcgaga gcgagagcga gagcgagagg ggaggacatg gaagcgaatg      300
ccgggatggg ggccggatcc tacaagcga aagagctggg ccggatacgc cagcactccg      360
acagcgcgcc caagcccctg aagcacttgg atggccagat gtgtcagatt tgtggtgata      420
ccgttggaact ttcggccagt ggtgatgtgt      450

```

<210> 128  
 <211> 302  
 <212> DNA  
 <213> Eucalyptus grandis

```

<400> 128
tcttgcccag gtttactcaa ggttttaggt ggagtcaaca caaacttcac cgtcacctca      60
aaagcagcgg atgatggagc attctcagaa ctctatatct ttaaaggac atcgctgttg      120
atccgcacca tgactctcct gatcatgaac attgtatgga ggctgttggc gggatctccg      180
atgccatcaa taatgggtat gattcggtgg gtccctctt tggtaggcta ttttcgcct      240
tctgggtcgc gtccatctct acccattcct aagggtatgc tcgggaagca agaccggatg      300
cc                                                                                   302

```

<210> 129  
 <211> 177  
 <212> PRT  
 <213> Eucalyptus grandis

```

<400> 129
Val Thr Gln Phe Leu Thr Asn Asn Asn Arg Ala Ser Gly Thr Leu Ser
 1          5          10          15
Ser Ile Arg Arg Phe Tyr Val Gln Asp Gly Lys Val Ile Pro Asn Ser
 20          25          30
Met Val Asn Leu Ser Gly Leu Pro Lys Val Asn Ser Ile Thr Ser Asp
 35          40          45
Tyr Cys Thr Ala Lys Met Asp Val Leu Asp Asp Ser Thr Ala Phe Asn
 50          55          60
Val His Gly Gly Leu Ala Lys Met Gly Lys Ser Leu Ala Arg Gly Ala
 65          70          75          80
Val Leu Val Val Ser Leu Trp Asp Asp Leu Gly Gly Gly Met Thr Trp
 85          90          95
Leu Asp Gly Leu Ala Gly Asp Ala Ser Ala Pro Gly Thr Leu Arg Gly
100          105          110
Pro Cys Thr Ala Ala Asn Val Thr Ser Asp Pro Ala Thr Ser Val Thr
115          120          125
Phe Ser Asn Ile Arg Val Gly Asp Ile Asn Ser Thr Phe Ser Gln Val
130          135          140
His Phe Gly Gln Cys Gly Gly Gln Phe Tyr Glu Gly Pro Ala Leu Cys
145          150          155          160
Ala Asp Pro Phe Glu Cys Val Phe Ser Asn Pro Tyr Tyr Ser Gln Cys
165          170          175
Leu

```

<210> 130  
 <211> 118  
 <212> PRT  
 <213> Eucalyptus grandis

```

<400> 130
Met Phe Asp Lys Ala Ala Leu Phe Ala Phe Ser Leu Leu Ala Val Thr
 1          5          10          15

```

Tyr Gly Gln Gln Val Gly Thr Gln Thr Ala Glu Ser His Pro Pro Leu  
                   20                  25                  30  
 Thr Trp Gln Lys Cys Thr Thr Ala Gly Gly Cys Thr Asn Val Ser Gly  
           35                  40                  45  
 Gly Ser Val Val Ile Asp Ala Asn Trp Arg Trp Val His Ser Ile Asn  
           50                  55                  60  
 Gly Thr Thr Asn Cys Tyr Thr Gly Gln Ala Trp Asn Thr Thr Leu Cys  
   65                  70                  75                  80  
 Pro Asp Asp Thr Thr Cys Ala Ala Asn Cys Ala Leu Asp Gly Ala Asp  
                   85                  90                  95  
 Tyr Ser Gly Thr Tyr Gly Ile Thr Thr Ser Gly Asn Ala Leu Thr Leu  
                   100                  105                  110  
 Lys Phe Val Thr Gln Ser  
           115

<210> 131  
 <211> 93  
 <212> PRT  
 <213> Eucalyptus grandis

<400> 131  
 Tyr Tyr Asp Ala Asn Leu Ile Ala Ile Ala Val Gly Phe Ser Arg Thr  
   1                  5                  10                  15  
 Ile Tyr Ser Val Ile Pro Gln Trp Ser Arg Leu Ile Gly Gly Val Phe  
           20                  25                  30  
 Phe Ser Phe Trp Val Leu Ala His Leu Tyr Pro Phe Ala Lys Gly Leu  
           35                  40                  45  
 Met Gly Arg Arg Gly Arg Thr Pro Thr Ile Val Phe Val Trp Ser Gly  
   50                  55                  60  
 Leu Ile Ala Ile Thr Ile Ser Leu Leu Trp Val Ala Ile Ser Pro Pro  
   65                  70                  75                  80  
 Ala Gly Ser Thr Gln Ile Gly Gly Ser Phe Gln Phe Pro  
                   85                  90

<210> 132  
 <211> 196  
 <212> PRT  
 <213> Eucalyptus grandis

<400> 132  
 Thr Thr Leu Glu Asp Gly Gly Val Pro Pro Asp Ala Ser Pro Ala Ser  
   1                  5                  10                  15  
 Leu Leu Lys Glu Ala Ile Gln Val Ile Ser Cys Gly Tyr Glu Asp Lys  
           20                  25                  30  
 Thr Glu Trp Gly Lys Glu Val Gly Trp Ile Tyr Gly Ser Val Thr Glu  
           35                  40                  45  
 Asp Ile Leu Thr Gly Phe Lys Met His Cys His Gly Trp Arg Ser Val  
   50                  55                  60  
 Tyr Cys Ile Pro Lys Arg Pro Ala Phe Lys Gly Ser Ala Pro Ile Asn  
   65                  70                  75                  80  
 Leu Ser Asp Arg Leu His Gln Val Leu Arg Trp Ala Leu Gly Ser Val  
                   85                  90                  95  
 Glu Ile Phe Leu Ser Arg His Cys Pro Ile Trp Tyr Gly Tyr Gly Gly  
                   100                  105                  110  
 Gly Leu Lys Trp Leu Glu Arg Phe Ser Tyr Ile Asn Ser Val Val Tyr  
                   115                  120                  125  
 Pro Trp Thr Ser Ile Pro Leu Ile Val Tyr Cys Ser Leu Pro Ala Ile

130                      135                      140  
 Cys Leu Leu Thr Gly Gln Phe Ile Val Pro Glu Ile Ser Asn Tyr Ala  
 145                      150                      155                      160  
 Ser Leu Val Phe Met Ala Leu Phe Ile Ser Ile Ala Ala Thr Gly Ile  
 165                      170                      175  
 Leu Glu Met Gln Trp Gly Gly Val Gly Ile Asp Asp Trp Trp Arg Asn  
 180                      185                      190  
 Glu Gln Phe Trp  
 195

<210> 133  
 <211> 196  
 <212> PRT  
 <213> Eucalyptus grandis

<400> 133  
 Thr Ala Leu Ile Pro Lys Lys Phe Gly Asn Ser Tyr Met Phe Ile Asp  
 1                      5                      10                      15  
 Ser Ile Pro Leu Ala Glu Phe Gln Gly Arg Pro Leu Ala Asp His Pro  
 20                      25                      30  
 Ser Val Lys Asn Gly Arg Pro Pro Gly Ala Leu Thr Val Leu Arg Arg  
 35                      40                      45  
 Leu Leu Asp Ala Ser Thr Val Ala Glu Ala Ile Ser Ala Ile Ser Cys  
 50                      55                      60  
 Trp Tyr Glu Asp Lys Thr Glu Trp Gly Glu Arg Val Gly Trp Ile Tyr  
 65                      70                      75                      80  
 Gly Ser Val Thr Glu Asp Val Val Thr Gly Tyr Arg Met His Asn Arg  
 85                      90                      95  
 Gly Trp Thr Ser Val Tyr Cys Val Thr Lys Arg Asp Ala Phe Arg Gly  
 100                      105                      110  
 Thr Ala Pro Ile Asn Leu Thr Asp Arg Leu His Gln Val Leu Arg Trp  
 115                      120                      125  
 Ala Thr Gly Ser Val Glu Ile Phe Phe Ser Arg Asn Asn Ala Ile Met  
 130                      135                      140  
 Ala Ser Gly Arg Leu Lys Phe Leu Gln Arg Ile Ala Tyr Leu Asn Val  
 145                      150                      155                      160  
 Gly Ile Tyr Pro Phe Thr Ser Ile Phe Leu Ile Val Tyr Cys Phe Leu  
 165                      170                      175  
 Pro Ala Leu Ser Leu Phe Ser Gly Lys Phe Ile Val Gln Ser Leu Ser  
 180                      185                      190  
 Val Ser Phe Leu  
 195

<210> 134  
 <211> 206  
 <212> PRT  
 <213> Eucalyptus grandis

<400> 134  
 Gly Lys Gly Gly Asp Lys Asn Tyr Ile Asp Lys Lys Arg Ala Gly Lys  
 1                      5                      10                      15  
 Arg Thr Glu Ser Asn Ile Pro Ile Phe Asn Met Glu Asp Ile Glu Glu  
 20                      25                      30  
 Gly Met Glu Gly Tyr Asp Asp Glu Arg Ser Leu Leu Met Ser Gln Lys  
 35                      40                      45  
 Ser Leu Glu Lys Arg Phe Gly Gln Ser Pro Val Phe Ile Ala Ala Thr  
 50                      55                      60



Phe Met Glu Gln Gly Gly Leu Pro Pro Ser Thr Asn Pro Ala Thr Leu  
 65 70 75 80  
 Leu Lys Glu Ala Ile His Val Ile Ser Cys Gly Tyr Glu Asp Lys Thr  
 85 90 95  
 Glu Trp Gly Lys Glu Ile Gly Trp Ile Tyr Gly Ser Val Thr Glu Asp  
 100 105 110  
 Ile Leu Thr Gly Phe Lys Met His Ala Arg Gly Trp Ile Ser Ile Tyr  
 115 120 125  
 Cys Met Pro Pro Arg Pro Ala Phe Lys Gly Ser Ala Pro Ile Asn Leu  
 130 135 140  
 Ser Asp Arg Leu Asn Gln Val Leu Arg Trp Ala Leu Gly Ser Ile Glu  
 145 150 155 160  
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<210> 135  
 <211> 126  
 <212> PRT  
 <213> Eucalyptus grandis

<400> 135  
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 <211> 51  
 <212> PRT  
 <213> Eucalyptus grandis

<400> 136  
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 <213> Eucalyptus grandis

<400> 137  
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 <213> Eucalyptus grandis

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&lt;210&gt; 140

&lt;211&gt; 1592

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 140

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&lt;210&gt; 141

&lt;211&gt; 3747

&lt;212&gt; DNA

<213> *Eucalyptus grandis*

&lt;400&gt; 141

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 <211> 770  
 <212> DNA  
 <213> *Eucalyptus grandis*

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<210> 143  
 <211> 543  
 <212> DNA  
 <213> *Pinus radiata*

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ggaaagacga gagtcattcc gctatcttga gatgttttac atcttgaagt acagaaatct 180

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accatattct cgtaacgcca tctcgttttc tcccatattt cttggtcctg gggttgctgt      420
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aaa

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&lt;210&gt; 144

&lt;211&gt; 805

&lt;212&gt; PRT

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 144

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Arg Val Glu Ala Lys Gly Lys Gly Ile Leu Gln Arg His Gln Ile Phe
35           40           45
Ala Glu Phe Glu Ala Ile Ser Glu Glu Ser Arg Ala Lys Leu Leu Asp
50           55           60
Gly Ala Phe Gly Glu Val Leu Lys Ser Thr Gln Glu Ala Ile Val Ser
65           70           75           80
Pro Pro Trp Val Ala Leu Ala Val Arg Pro Arg Pro Gly Val Trp Glu
85           90           95
His Ile Arg Val Asn Val His Ala Leu Val Leu Glu Gln Leu Glu Val
100          105          110
Ala Glu Tyr Leu His Phe Lys Glu Glu Leu Ala Asp Gly Ser Leu Asn
115          120          125
Gly Asn Phe Val Leu Glu Leu Asp Phe Glu Pro Phe Thr Ala Ser Phe
130          135          140
Pro Arg Pro Thr Leu Ser Lys Ser Ile Gly Asn Gly Val Glu Phe Leu
145          150          155          160
Asn Arg His Leu Ser Ala Lys Leu Phe His Asp Lys Glu Ser Leu His
165          170          175
Pro Leu Leu Glu Phe Leu Gln Val His Cys Tyr Lys Gly Lys Asn Met
180          185          190
Met Val Asn Ala Arg Ile Gln Asn Val Phe Ser Leu Gln His Val Leu
195          200          205
Arg Lys Ala Glu Glu Tyr Leu Thr Ser Leu Lys Pro Glu Thr Pro Tyr
210          215          220
Ser Gln Phe Glu His Lys Phe Gln Glu Ile Gly Leu Glu Arg Gly Trp
225          230          235          240
Gly Asp Thr Ala Glu Arg Val Leu Glu Met Ile Gln Leu Leu Leu Asp
245          250          255
Leu Leu Glu Ala Pro Asp Pro Cys Thr Leu Glu Lys Phe Leu Asp Arg
260          265          270
Val Pro Met Val Phe Asn Val Val Ile Met Ser Pro His Gly Tyr Phe

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385	390	395
Glu Leu Gln Gly Lys Pro Asp Leu Ile Ile Gly Asn Tyr Ser Asp Gly		
405	410	415
Asn Ile Val Ala Ser Leu Leu Ala His Lys Leu Gly Val Thr Gln Cys		
420	425	430
Thr Ile Ala His Ala Leu Glu Lys Thr Lys Tyr Pro Glu Ser Asp Ile		
435	440	445
Tyr Trp Lys Lys Phe Glu Glu Lys Tyr His Phe Ser Cys Gln Phe Thr		
450	455	460
Ala Asp Leu Ile Ala Met Asn His Thr Asp Phe Ile Ile Thr Ser Thr		
465	470	475
Phe Gln Glu Ile Ala Gly Ser Lys Asp Thr Val Gly Gln Tyr Glu Ser		
485	490	495
His Met Asn Phe Thr Leu Pro Gly Leu Tyr Arg Val Val His Gly Ile		
500	505	510
Asp Val Phe Asp Pro Lys Phe Asn Ile Val Ser Pro Gly Ala Asp Met		
515	520	525
Ser Ile Tyr Phe Ala Tyr Thr Glu Gln Glu Arg Arg Leu Lys Ser Phe		
530	535	540
His Pro Glu Ile Glu Glu Leu Leu Phe Ser Asp Val Glu Asn Lys Glu		
545	550	555
His Leu Cys Val Leu Lys Asp Lys Lys Lys Pro Ile Ile Phe Thr Met		
565	570	575
Ala Arg Leu Asp Arg Val Lys Asn Leu Thr Gly Leu Val Glu Trp Tyr		
580	585	590
Gly Lys Asn Ser Lys Leu Arg Glu Leu Ala Asn Leu Val Val Val Gly		
595	600	605
Gly Asp Arg Arg Lys Asp Ser Lys Asp Leu Glu Glu Gln Ser Glu Met		
610	615	620
Lys Lys Met Tyr Asp Leu Ile Glu Lys Tyr Lys Leu Asn Gly Gln Phe		
625	630	635
Arg Trp Ile Ser Ser Gln Met Asn Arg Val Arg Asn Gly Glu Leu Tyr		
645	650	655
Arg Tyr Ile Cys Asp Thr Lys Gly Val Phe Val Gln Pro Ala Ile Tyr		
660	665	670
Glu Ala Phe Gly Leu Thr Val Val Glu Ala Met Thr Cys Gly Leu Pro		
675	680	685
Thr Phe Ala Thr Cys Asn Gly Gly Pro Ala Glu Ile Ile Val His Gly		
690	695	700
Lys Ser Gly Tyr His Ile Asp Pro Tyr His Gly Asp Gln Ala Ala Glu		
705	710	715
Leu Leu Val Asp Phe Phe Asn Lys Cys Lys Ile Asp Gln Ser His Trp		
725	730	735

Asp Glu Ile Ser Lys Gly Ala Met Gln Arg Ile Glu Glu Lys Tyr Thr  
                   740                  745                  750  
 Trp Lys Ile Tyr Ser Glu Arg Leu Leu Asn Leu Thr Ala Val Tyr Gly  
           755                  760                  765  
 Phe Trp Lys His Val Thr Asn Leu Asp Arg Arg Glu Ser Arg Arg Tyr  
       770                  775                  780  
 Leu Glu Met Phe Tyr Ala Leu Lys Tyr Arg Pro Leu Ala Gln Ser Val  
 785                  790                  795                  800  
 Pro Pro Ala Val Glu  
                   805

<210> 145  
 <211> 419  
 <212> PRT  
 <213> Pinus radiata

<400> 145  
 Met Gly Tyr Pro Val Gln Glu Val Ser Lys Glu His Asp Gln Trp Ala  
   1                  5                  10                  15  
 Gly Phe Val Glu Gly Glu Ser Val Leu Gln Arg Gly Arg Glu Ile Leu  
           20                  25                  30  
 Leu Gln Gly Phe Asn Trp Glu Ser His Lys Tyr Lys Trp Trp Pro Asn  
       35                  40                  45  
 Leu Glu Glu Lys Ile Pro His Ile Ala Lys Ala Gly Phe Thr Ser Val  
       50                  55                  60  
 Trp Leu Pro Pro Ala Phe Asp Ser Ala Ala Pro Gln Gly Tyr Leu Pro  
 65                  70                  75                  80  
 Arg Asn Ile Tyr Ser Leu Asn Ser Ala Tyr Gly Ser Glu Tyr Gln Leu  
                   85                  90                  95  
 Lys Ser Leu Leu Met Thr Met Arg Lys Lys Asn Val Arg Ala Met Ala  
           100                  105                  110  
 Asp Ile Val Ile Asn His Arg Met Gly Ser Ser Gln Gly Phe Gly Gly  
       115                  120                  125  
 Leu Tyr Asn Arg Tyr Asp Gly Leu Pro Leu Pro Trp Asp Glu Arg Ala  
       130                  135                  140  
 Val Thr Arg Cys Ser Gly Gly Leu Gly Asn Trp Ser Thr Gly Asp Asn  
 145                  150                  155                  160  
 Phe His Gly Val Pro Asn Val Asp His Thr Gln Asp Phe Val Arg Lys  
                   165                  170                  175  
 Asp Ile Lys Asp Trp Leu Arg Trp Leu Arg Lys Ser Val Gly Phe Gln  
           180                  185                  190  
 Asp Phe Arg Phe Asp Phe Ala Lys Gly Tyr Ala Ala Lys Phe Val Lys  
       195                  200                  205  
 Glu Tyr Ile Glu Ala Ser Thr Pro Met Phe Ala Val Gly Glu Tyr Trp  
       210                  215                  220  
 Asp Ser Cys Asn Tyr Thr Pro Pro Ser Tyr His Leu Asp Lys Asn Gln  
 225                  230                  235                  240  
 Asp Ser His Arg Gln Arg Ile Ile Asn Trp Ile Asp Gly Thr Ser Gly  
                   245                  250                  255  
 Ile Ser Ala Ala Phe Asp Phe Thr Thr Lys Gly Ile Leu Gln Glu Ala  
           260                  265                  270  
 Val Lys Gly Gln Cys Trp Arg Leu Arg Asp His Gln Gly Lys Pro Pro  
       275                  280                  285  
 Gly Val Leu Gly Trp Trp Pro Pro Arg Leu Val Leu Leu Asn Glu Asn  
       290                  295                  300  
 His Asp Thr Gly Ser Thr Gln Gly His Trp Pro Phe Pro Cys Asp His  
 305                  310                  315                  320



[illegible]

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<210> 146
<211> 955
<212> PRT
<213> Eucalyptus grandis
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<400> 146															
Met	Met	Glu	Ser	Gly	Val	Pro	Leu	Cys	Asn	Thr	Cys	Gly	Glu	Ala	Val
1				5					10					15	
Gly	Val	Asp	Glu	Lys	Gly	Glu	Val	Phe	Val	Ala	Cys	Gln	Glu	Cys	Asn
			20					25					30		
Phe	Ala	Ile	Cys	Lys	Ala	Cys	Val	Glu	Tyr	Glu	Ile	Lys	Glu	Gly	Arg
		35					40					45			
Lys	Ala	Cys	Leu	Arg	Cys	Gly	Thr	Pro	Phe	Glu	Ala	Asn	Ser	Met	Ala
	50					55					60				
Asp	Ala	Glu	Arg	Asn	Glu	Leu	Gly	Ser	Arg	Ser	Thr	Met	Ala	Ala	Gln
65					70					75				80	
Leu	Asn	Asp	Pro	Gln	Asp	Thr	Gly	Ile	His	Ala	Arg	His	Ile	Ser	Ser
				85					90					95	
Val	Ser	Thr	Leu	Asp	Ser	Glu	Tyr	Asn	Asp	Glu	Thr	Gly	Asn	Pro	Ile
			100					105					110		
Trp	Lys	Asn	Arg	Val	Glu	Ser	Trp	Lys	Asp	Lys	Lys	Asn	Lys	Lys	Lys
		115					120					125			
Lys	Ala	Pro	Thr	Lys	Ala	Glu	Lys	Glu	Ala	Gln	Val	Pro	Pro	Glu	Gln
		130				135					140				
Gln	Met	Glu	Glu	Lys	Gln	Ile	Ala	Asp	Ala	Ser	Glu	Pro	Leu	Ser	Thr
145					150				155					160	
Val	Ile	Pro	Ile	Ala	Lys	Ser	Lys	Leu	Ala	Pro	Tyr	Arg	Thr	Val	Ile
			165					170						175	
Ile	Met	Arg	Leu	Ile	Ile	Leu	Ala	Leu	Phe	Phe	His	Tyr	Arg	Val	Thr
			180					185					190		
His	Pro	Val	Asp	Ser	Ala	Tyr	Pro	Leu	Trp	Leu	Thr	Ser	Ile	Ile	Cys
		195					200					205			
Glu	Ile	Trp	Phe	Ala	Tyr	Ser	Trp	Val	Leu	Asp	Gln	Phe	Pro	Lys	Trp
		210				215					220				
Ser	Pro	Val	Asn	Arg	Ile	Thr	His	Val	Asp	Arg	Leu	Ser	Ala	Arg	Tyr
225					230					235				240	
Glu	Lys	Glu	Gly	Glu	Pro	Ser	Glu	Leu	Ala	Ala	Val	Asp	Phe	Phe	Val
			245						250					255	
Ser	Thr	Val	Asp	Pro	Met	Lys	Glu	Pro	Pro	Leu	Ile	Thr	Ala	Asn	Thr
			260					265					270		
Val	Leu	Ser	Ile	Leu	Ala	Val	Asp	Tyr	Pro	Val	Asp	Lys	Val	Ser	Cys
		275					280					285			

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Tyr Leu Ser Asp Asp Gly Ala Ala Met Leu Ser Phe Glu Ser Leu Val
290                295                300
Glu Thr Ala Asp Phe Ala Arg Lys Trp Val Pro Phe Cys Lys Lys Tyr
305                310                315                320
Ser Ile Glu Pro Arg Ala Pro Glu Phe Tyr Phe Ser Gln Lys Ile Asp
325                330                335
Tyr Leu Lys Asp Lys Ile Gln Pro Ser Phe Val Lys Glu Arg Arg Ala
340                345                350
Met Lys Arg Asp Tyr Glu Glu Phe Lys Val Arg Val Asn Ala Leu Val
355                360                365
Ala Lys Ala Gln Lys Ala Pro Glu Glu Gly Trp Ser Met Gln Asp Gly
370                375                380
Thr Pro Trp Pro Gly Asn Asn Ser Arg Asp His Pro Gly Met Ile Gln
385                390                395                400
Val Phe Leu Gly Ser Ser Gly Ala His Asp Ile Glu Gly Asn Glu Leu
405                410                415
Pro Arg Leu Val Tyr Val Ser Arg Glu Lys Arg Pro Gly Phe Gln His
420                425                430
His Lys Lys Ala Gly Ala Glu Asn Ala Leu Val Arg Val Ser Ala Ile
435                440                445
Leu Thr Asn Ala Pro Tyr Ile Leu Asn Leu Asp Cys Asp His Tyr Val
450                455                460
Asn Tyr Ser Asn Ala Val Arg Glu Ala Met Cys Phe Leu Met Asp Pro
465                470                475                480
Gln Val Gly Arg Asn Leu Cys Tyr Val Gln Phe Pro Gln Arg Phe Asp
485                490                495
Gly Ile Asp Arg Ser Asp Arg Tyr Ala Asn Arg Asn Thr Val Phe Phe
500                505                510
Asp Val Asn Met Lys Gly Leu Asp Gly Ile Gln Gly Pro Val Tyr Val
515                520                525
Gly Thr Gly Cys Val Phe Asn Arg Gln Ala Leu Tyr Gly Tyr Gly Pro
530                535                540
Pro Ser Met Pro Asn Leu Pro Lys Pro Ser Ser Ser Cys Ser Trp Cys
545                550                555                560
Gly Cys Cys Ser Cys Cys Cys Pro Ser Lys Lys Pro Thr Lys Asp Leu
565                570                575
Ser Glu Val Tyr Arg Asp Ser Lys Arg Glu Asp Leu Asn Ala Ala Ile
580                585                590
Phe Asn Leu Gly Glu Ile Asp Asn Tyr Asp Glu His Glu Arg Ser Met
595                600                605
Leu Ile Ser Gln Met Ser Phe Glu Lys Thr Phe Gly Leu Ser Thr Val
610                615                620
Phe Ile Glu Ser Thr Leu Leu Ala Asn Gly Gly Val Pro Glu Ser Ala
625                630                635                640
His Pro Ser Met Leu Ile Lys Glu Ala Ile His Val Ile Ser Cys Gly
645                650                655
Tyr Glu Glu Lys Thr Ala Trp Gly Lys Glu Ile Gly Trp Ile Tyr Gly
660                665                670
Ser Val Thr Glu Asp Ile Leu Thr Gly Phe Lys Met His Cys Arg Gly
675                680                685
Trp Arg Ser Val Tyr Cys Met Pro Leu Arg Pro Ala Phe Lys Gly Ser
690                695                700
Ala Pro Ile Asn Leu Ser Asp Arg Leu His Gln Val Leu Arg Trp Ala
705                710                715                720
Leu Gly Ser Val Glu Ile Phe Leu Ser Arg His Cys Pro Leu Trp Tyr
725                730                735
Gly Phe Gly Gly Gly Arg Leu Lys Trp Leu Gln Arg Leu Ala Tyr Ile

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      740      745      750
Asn Thr Ile Val Tyr Pro Phe Thr Ser Leu Pro Leu Val Ala Tyr Cys
      755      760      765
Thr Ile Pro Ala Ile Cys Leu Thr Gly Lys Phe Ile Ile Pro Thr
      770      775      780
Leu Ser Asn Leu Ala Ser Val Leu Phe Leu Gly Leu Phe Leu Ser Ile
785      790      795      800
Ile Val Thr Ser Val Leu Glu Leu Arg Trp Ser Gly Val Ser Ile Glu
      805      810      815
Asp Trp Trp Arg Asn Glu Gln Phe Trp Val Ile Gly Gly Val Ser Ala
      820      825      830
His Leu Phe Ala Val Phe Gln Gly Phe Leu Lys Met Leu Ala Gly Leu
      835      840      845
Asp Thr Asn Phe Thr Val Thr Thr Lys Ala Ala Asp Asp Ala Glu Phe
      850      855      860
Gly Glu Leu Tyr Met Ile Lys Trp Thr Thr Leu Leu Ile Pro Pro Thr
865      870      875      880
Thr Leu Leu Ile Val Asn Met Val Gly Val Val Ala Gly Phe Ser Asp
      885      890      895
Ala Leu Asn Lys Gly Tyr Glu Ala Trp Gly Pro Leu Phe Gly Lys Val
      900      905      910
Phe Phe Ala Phe Trp Val Ile Leu His Leu Tyr Pro Phe Leu Lys Gly
      915      920      925
Leu Met Gly Arg Gln Asn Arg Thr Pro Thr Ile Val Val Leu Trp Ser
      930      935      940
Val Phe Trp Leu Leu Ser Ser Leu Ser Ser Gly
945      950      955

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<210> 147  
 <211> 124  
 <212> PRT  
 <213> *Eucalyptus grandis*

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      <400> 147
Leu Lys Asp Ala Ile Ile Ser His Gly Cys Phe Leu Arg Glu Cys Arg
 1      5      10      15
Val Glu Arg Ser Ile Val Gly Glu Arg Ser Arg Leu Asp Ser Gly Val
      20      25      30
Glu Leu Lys Asp Thr Val Met Met Gly Ala Asp Tyr Tyr Gln Thr Glu
      35      40      45
Ser Glu Ile Ala Ser Leu Leu Ala Glu Gly Lys Val Pro Ile Gly Ile
      50      55      60
Gly Lys Asn Thr Lys Ile Arg Asn Cys Ile Ile Asp Lys Asn Ala Lys
65      70      75      80
Ile Gly Lys Asp Val Ala Ile Val Asn Lys Asp Gly Val Glu Glu Ala
      85      90      95
Asp Arg Pro Gly Asp Gly Phe Tyr Ile Arg Leu Gly Ile Thr Val Ile
      100      105      110
Leu Glu Lys Ala Thr Ile Glu Asp Gly Thr Val Ile
      115      120

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<210> 148  
 <211> 80  
 <212> PRT  
 <213> *Pinus radiata*

<400> 148

Asp Thr Ile Ser Asn Gly Gly Leu Gln Arg Ile Tyr Glu Arg Tyr Thr  
 1 5 10 15  
 Trp Lys Ile Tyr Ala Glu Lys Leu Met Thr Leu Ala Gly Val Tyr Gly  
 20 25 30  
 Phe Trp Lys Tyr Val Ser Lys Leu Glu Arg Arg Glu Ser Phe Arg Tyr  
 35 40 45  
 Leu Glu Met Phe Tyr Ile Leu Lys Tyr Arg Asn Leu Val Lys Thr Val  
 50 55 60  
 Pro Phe Ser Val Glu Glu Ser Thr Asp Gly Ile Glu Glu Lys Ser Ala  
 65 70 75 80

&lt;210&gt; 149

&lt;211&gt; 375

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 149

atccgctggg	agggttagga	ctgacatcag	cgatctcatg	acccaaattg	ttgagttcct	60
agccaagaac	agcgcacctc	tcaccgtcaa	catttatecca	ttcctgagtc	tctatggtaa	120
cgacaacttc	cccatcgact	atgccttctt	cgatggggcc	acccagtcg	tggacaacgg	180
gatacaatac	acgaacgtgt	tcgatgccaa	cttcgacact	ctagtatcgg	ctcttaaggc	240
gggtgggcat	ggggacatga	ccctcatggt	gggtgaagtg	ggatggccta	cagatgggtga	300
caagaatgcc	aatatagcta	gtgctgtag	attctacaac	gggtcttatgc	cgaggctcgc	360
ggccaatact	gggac					375

&lt;210&gt; 150

&lt;211&gt; 356

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 150

gaacaacaac	tctccattgc	tagccaacat	ctacacctac	ttcagctacg	ttggtaacct	60
gaaagacatc	agcctgccct	acgccctatt	cacctcgccg	tccgctcgcg	tccgagacgg	120
agcccatgag	taccggaatc	tggtcgacgc	gatgctggac	gccctctact	ccgcactcga	180
gagggtgggc	ggggctgccc	tcaggggtgt	gggtcggag	agcggctggc	catccgcggg	240
cgcgctcgc	gcgacggctc	acaacgcgag	gacatacaac	gggaatctga	tcaagcacgt	300
gaagggcggc	acgcgaagag	gccgaacggg	gcgatcgaga	cctacatatt	cgcctt	356

&lt;210&gt; 151

&lt;211&gt; 470

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 151

cttcctcagg	gagaccggct	cgcttttctt	gggtgaacctg	tacctatttt	tccgcttctc	60
caaggacact	ctcgattacg	cgctgttcag	gccgaatgca	gggggtgatg	atgagaattc	120
caaacttggt	tacactaaca	tggtggacgg	gcaattggac	gctgtctact	cggcgatgaa	180
gggtgctggc	ttaccgata	tcgagatcgt	gatagccgaa	acaggatggc	cctcattgtg	240
tgactcgacc	caagtcggcg	tggtatgcaa	gacggcggca	gaatacaaca	gtaattctcat	300
ccgccatgta	tcgtcgggcg	ccggcacgcc	cctcatgcca	aaacggacat	ttgagacctt	360

catttttgcct ctcttcaacg agaatttgaa gccgggaccg acgtgcgaga ggaacttcgg 420  
actcttcagg ccggacatga caccggtgta tgatgccggg atcttgaggc 470

<210> 152

<211> 412

<212> DNA

<213> Eucalyptus grandis

<400> 152

tgattttattg tgattgccat tctctcttctt tcttgattct gcatgttaca cgggtgcagca 60  
actctcttat tttaacatca ctgggcaggt gtgcccttat tccttaatac aaggcagcgt 120  
gcaccattta gttgaaatta atagggtatt ttggctgtct attgggcgaa aagaggcccc 180  
aatgtttctc gtactccggg ttcttcttgt tctcgtcgaa catggcaaat atgtaagttt 240  
cgatcggttt tccgggcttc ttgggagtcc cttgcttcac gtgctgaatc aaattcgagt 300  
tgtaaattct cgcgttttca atggacgttc ccttcccgcc agcagatggg ccaaccactc 360  
tccgacacaa ccacttctag ggaccccccg ccggatttct ctagtgcaga gt 412

<210> 153

<211> 328

<212> DNA

<213> Eucalyptus grandis

<400> 153

gaacaacaac tctccattgc tagccaacat ctacacctac ttcagctacg ttggttaacc 60  
gaaagacatc agcctgccct acgccctatt cacctcgccg tcggtcgtcg tccgagacgg 120  
agcccatgag taccggaatc tggtcgacgc gatgtggac gccctctact ccgcactcga 180  
gagggtggc ggggtgccc tcagggtggg ggtctcggag agcggctggc catccgcggg 240  
cgcgttcgct gcgacggtcg acaacgcgag gacatacaac gggaaatctga tcaagcacgt 300  
gaaggcgccg acgccgaaga ggccgaac 328

<210> 154

<211> 373

<212> DNA

<213> Eucalyptus grandis

<400> 154

gttaacgcct accccttctt cgcctaagag tccaactccg acgtcatctc cctggactac 60  
gccctcttcc gggagaaccc gggcgtcgtc gacgccggca acgggctccg ctacttcaac 120  
ctcttcgacg cccagatcga cgcctcttcc gccgccatgt cggccctcaa gtacgacgac 180  
atcaagatgg tcgtcaccga gacgggctgg ccctccaagg gcgacgagaa cgaggctcggc 240  
gccagcaagg acaatgccgc cgcctacaac ggcaacctcg tccgccggat cctcaccggc 300  
ggcggcaccc ctctgaggcg gcaggccgac ctaccgtct acctcttcgc gctcttcaac 360  
gagaacaaga agc 373

<210> 155

<211> 465

<212> DNA

<213> Eucalyptus grandis

<400> 155

cgatcatcag tttcttggtg agcaacaact ctccattgct agccaacatc tacacctact 60  
tcagctacgt cgtaaacccg aaagacatca gcctgcccta cgccctattc acctcgccat 120  
cggctcgtct ccgagacggg gccatgagt accggaacct gttcgacgag atgctggacg 180  
ccctctactc cgcactcgag agggctggcg gggctgccct ccgggtgggt gtctcggaga 240  
gcggctggcc atccgcgggc gcgttcgctg cgacggctga caacgcgagg acatacaacg 300  
ggaatctgat caagcacgtg aagggcggca gcgcgaagag gccgaacggg gcgatcgaga 360  
cctacatatt cgccttgctc gacgagaacc agaagcagcc ggagctggag aagcacttcg 420

ggctcttctt ccccaacaag cagcccaagt acccgctcag ctttg 465

<210> 156  
<211> 359  
<212> DNA  
<213> Eucalyptus grandis

<400> 156

ccagatcaag gtctcaacag ctgtggacac ggcgcatttta ggagaatcgt ctctccatc	60
taagggcaca cttaaggccg atcacagacc tctcctagat cctataatca cattcctagt	120
caacaacaag tcccctctgc ttgtcaacat ctatccgtac ttcagctaca gcgacaatcc	180
caacgaacgc ctcgactatg ctctgttcac ggcgaactcg gttgtggtgt cggatggagc	240
acttgggtac cggaacttgt ttgacgcaat tctagatgct gtttactctg cactagagaa	300
atccggcggg ggggccctag aagtgggtgt gtcggagagt ggttgccat ctgctggcg	359

<210> 157  
<211> 325  
<212> DNA  
<213> Eucalyptus grandis

<400> 157

ccagatcaag gtctcaacag ctgtggacac ggcgcatttta ggagaatcgt ctctccatc	60
taagggcaca cttaaggccg atcacagacc tctcctagat cctataatca cattcctagt	120
caacaacaag tcccctctgc ttgtcaacat ctatccgtac ttcagctaca gcgacaatcc	180
caacgaacgc ctcgactatg ctctgttcac ggcgaactcg gttgtggtgt cggatggagc	240
acttgggtac cggaacttgt ttgacgcaat tctagatgct gtttactctg cactagagaa	300
atccggcggg ggggccctag aagt	325

<210> 158  
<211> 362  
<212> DNA  
<213> Eucalyptus grandis

<400> 158

gtttgctatg ggatgctcgg gaacaacctg ccgtccgcgt cggaagtcgt cgccctctac	60
aagtcgccg gcatcaagca gatgagactc tacgacccta gccaaaccagc tctacaagcc	120
ctgagaggct cgaacatcga gctcatctc ggcgtcccca actcggagct ccaggccctt	180
gcttccaacc ccgccaacgc gaactcgtgg gtgcagagga acgtgaagaa ctactcggcc	240
ggcgtcaggt tccgctacat cgcgtgggc aacgaggtga gccctgtcaa cggaggcact	300
gcacaattcg ccaggttcgt cctccccgcg atgaggaaca tacaggccgc gctcagatcg	360
tc	362

<210> 159  
<211> 432  
<212> DNA  
<213> Eucalyptus grandis

<400> 159

taaatgccga tgtatacgaa tcaccagaag acaaccact tccatccgcc ggaacgttcc	60
gaagcgacat aagtgatgtc atgacccaaa tcgtcaagtt catggctgag aacaatgcac	120
ccttcaccgt gaacatttac ccgtttctga gtctttacgg caataatgac ttccctttca	180
attatgcttt ctctgacggg gcaactccaa ttgttgacaa ggggattgag tacaccaatg	240
tctttgatgc caactttgac actttggtgt cggctcttaa agcagttgga catgggaaca	300
tgaccatcat cataggcgag gtgggttggc ccacagaagg tgacataaat gcaaacaacg	360
gtaacgcgta caggttttat aatgggctct ttacaaaact tggagcgaat agagggactc	420
cacttcggcc ag	432

<210> 160  
 <211> 379  
 <212> DNA  
 <213> *Eucalyptus grandis*

<400> 160  
 caccgtacct gtgaacgcag acgtgtacaa ctcccccgtc agcaatcccg taccatccgc 60  
 tgggaggttt aggactgaca tcagcgatct catgacccaa attgtcgagt tcctagccaa 120  
 gaacagcgca cctctcaccg tcaacattta tccattcctg agtctctatg gtaacgacaa 180  
 cttcccatc gactatgcct tcttcgatgg ggccacccca gtcgtggaca acgggataca 240  
 atacacgaac gtgttcgatg ccaacttcga cactctagta tcggctctta aggtgggtggg 300  
 ccatggggac atgaccctca tgggtgggtga agtgggatgg cctacagatg gtgacaagaa 360  
 tgccaatata gctagtgc 379

<210> 161  
 <211> 361  
 <212> DNA  
 <213> *Eucalyptus grandis*

<400> 161  
 gtttgctatg ggatgctcgg gaacaacctg ccgtccgcgt cggaagtcgt cgccctctac 60  
 aagtcgccg gcatacgcga gatgagactc tacgacctta gcccaaccagc tctacaagcc 120  
 ctgagaggct cgaacatcga gctcatcctc ggcgccccca actcggagct ccaggccctt 180  
 gcttccaacc ccgccaacgc gaactcgtgg gtgcagagga acgtgaagaa ctactcgccc 240  
 ggcgtcaggt tccgctacat cgcctgtggc aacgaggtga gccctgtcaa cggaggcact 300  
 gcacaattcg ccaggttcgt cctccccgcg atgaggaaca tacaggccgc gctcagatcg 360  
 t 361

<210> 162  
 <211> 402  
 <212> DNA  
 <213> *Pinus radiata*

<400> 162  
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 gctcgggtgcc acgcccagaga atgcaaagac ttacaatggg aaccttttgg agaggctcca 240  
 caagaaggag ggtactcccc tgaagcccaa tgtgagcgtg caggccttca tttttgcgct 300  
 ctttaatgag aatttgaagt ccgggcctac atccgagaga aattatgggc tctttaaacc 360  
 agacggaacc gagacgtatg accttggctt gaaagggatt ga 402

<210> 163  
 <211> 297  
 <212> DNA  
 <213> *Pinus radiata*

<400> 163  
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 gcgtttataa caacaatctc atccgccatg tgcctcctaaa tgtagggact cccaagaggc 180  
 cgggaacgag cattgagacc tacatcttcg cacttttcaa cgagaacaga aaagctgggtg 240  
 atgagacgga gcgtcacttt gggcttttct accctaacca acaatttgta tactctg 297

<210> 164  
 <211> 427  
 <212> DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 164

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acaatgggaa	caggggttcc	atttctctgg	actacgctct	gtttagggtca	acctctaccg	180
tggtgcagga	cgagggtcgc	agctacatca	acttattcga	tgccctcgtc	gatacccttc	240
tttctgccat	ggaggacttg	gggtatcgca	acatcccact	catcgttact	gaaagcggat	300
ggccttctgg	tggcaatgat	gtggccacgg	ttgacaacgc	tcgcgtttat	aacaacaatc	360
tcatccgcca	tgtgctctca	aatgtaggga	ctcccaagag	gccgggaacg	agcattgaga	420
cctacat						427

&lt;210&gt; 165

&lt;211&gt; 205

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 165

ggttgacaac	gctcgcgttt	ataacaacaa	tctcatccgc	catgtgctct	caaagttagg	60
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cagaaaagct	ggatgatgaga	cggagcgtca	ctttgggctt	ttctacccta	accaacaatc	180
tgtatactct	ctaaacttta	ctccg				205

&lt;210&gt; 166

&lt;211&gt; 393

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 166

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tgtatactct	ctaaacttta	ctccgtaact	gcg			393

&lt;210&gt; 167

&lt;211&gt; 394

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 167

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&lt;210&gt; 168

&lt;211&gt; 498

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 168



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gcagctcata	tgcttcta					498

&lt;210&gt; 169

&lt;211&gt; 278

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 169

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actcccaaga	ggccgggaac	gagcattgag	acctcatc			278

&lt;210&gt; 170

&lt;211&gt; 419

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 170

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tgtatactct	ctaaacttta	ctccgtaact	gcgtcgcagt	ccgacgaacg	aatagagcc	419

&lt;210&gt; 171

&lt;211&gt; 437

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 171

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attcctgtca	gatcacgggt	ctcccttcat	ggctaacatc	tatccatact	tcagctacaa	240
tgggaacagg	ggttccattt	ctctggacta	cgctctgttt	aggtcaacct	ctaccgtggg	300
gcaggacgag	ggtcgcagct	acatcaactt	attcgatgcc	ctcgtcgata	cccttctttc	360
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&lt;210&gt; 172

&lt;211&gt; 343

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 172

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cccaccgtct	aaaggtgtct	tcaggaacna	ggttaaagat	atcatgagtt	ctctacttca	180
attcctgtca	gatcacgggt	ctcccctcat	ggctaacatc	tatccatact	tcagctacaa	240
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&lt;210&gt; 173

&lt;211&gt; 563

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 173

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accaacctgt	tcgatgccat	ggttgacact	cttttgtcgg	ccatggaagc	ctcgggggtat	120
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ggaacaccaa	agaggccagg	aatgagcatc	gacacatatg	ttttcgcgct	tttcaacgag	300
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aatgcaggca	gagttccctg	agt				563

&lt;210&gt; 174

&lt;211&gt; 639

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 174

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gcagaacgcc	aatctgcaga	ataacatcaa	ggctcgcacc	actcactcct	ccgatgttag	240
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aatctcatcc	gccatgtctt	atcgaatgaa	gggacacca			639

&lt;210&gt; 175

&lt;211&gt; 534

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 175

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gatgccatgg	tcgattctat	tttctccgca	atgggaagcct	tgggatattc	caacatccca	480
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<210> 176  
 <211> 345  
 <212> DNA  
 <213> Pinus radiata

<400> 176  
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 cattcaaaact gcgatccaga acgctaaact gcaggatagc atcaagggtct ctaccacca 180  
 caggccggat gtttagcagcg gctaccgcc gtctaaagga gtcttcgtag atgctgtgaa 240  
 ggacacgatg ggccaaatac tcaaatttct gtcacagaac ggcggtccct tcatggcgga 300  
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<210> 177  
 <211> 339  
 <212> DNA  
 <213> Pinus radiata

<400> 177  
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 aaagaggcca ggaatgagca tcgacacata tgttttcgag cttttcaacg aggatttgaa 180  
 gcaaggcgac gagacataga aacactttgg actattcgac cctaatacta aacagcctgt 240  
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 cgatgcagac aagcccttat cttcatgcac tgctatatg 339

<210> 178  
 <211> 313  
 <212> DNA  
 <213> Pinus radiata

<400> 178  
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 cctggacagc acattgatac atacattttc gctcttttca acgagaattt gaaaggcggg 180  
 gacgagccag aacgacattt tggacttttc tatectgac aaaaccttgt ttaccctgtt 240  
 aacttctccc cttaaattca tctgatctgt gttttgcatt agaatttcca gtattacca 300  
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<210> 179  
 <211> 460  
 <212> DNA  
 <213> Pinus radiata

<400> 179  
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 tcaaacgcga gtccagaacg ccaatctgca gagctccata aaggtctcca ctaccacgc 180  
 cacgtctgtt ctgggaaact cgtatcccc ttctcagga gaattcgctg atgaattgaa 240  
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 cgtgtatccc tacttcagtt acatttaca ccaggcccaa atctcgtag actatgcttt 360  
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<210> 180  
 <211> 296

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 180

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cctgttcgat	gcatgggttg	atactctttt	gtcggccatg	gaagcctctg	ggtatcccaa	180
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tgagaatgct	cagacctata	acaataatct	tattaaacat	gtattatcga	atgcag	296

&lt;210&gt; 181

&lt;211&gt; 351

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 181

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gacggtgatc	acagctacac	caacctgttc	gatgccatgg	gtgatactct	t	351

&lt;210&gt; 182

&lt;211&gt; 457

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 182

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&lt;210&gt; 183

&lt;211&gt; 358

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 183

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tgcaagcttt	ggcgaattct	ggtatcgatg	taatagtggg	tgctcgtaac	agcgaactgg	120
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&lt;210&gt; 184

&lt;211&gt; 348

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 184

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ccaggatagc	atcaaggctc	ctaccaccca	caggccggat	gtagcagcg	gctaccgcgc	120
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cacaaaagac	atttctctgg	attatgctct	gtttaagtct	acgtctacgg	tggtgcaaga	300
cgggtgatcac	agctacacca	acctgttcga	tgccatgggt	gatactct		348

&lt;210&gt; 185

&lt;211&gt; 594

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 185

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gttcagtagc	tacctatctc	tggtccggcc	atgaagaaca	ttcaaactgc	gatccagaac	300
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atcggcaaca	caaaagacat	ttctctggat	tatgctctgt	ttaagtctac	gtctacgggg	540
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&lt;210&gt; 186

&lt;211&gt; 360

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 186

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&lt;210&gt; 187

&lt;211&gt; 397

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 187

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ccctcttcag	agattttaca	gactctgccg	atcaggg			397

&lt;210&gt; 188

&lt;211&gt; 531

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 188

ggaagacaaa	gagaagagct	atattttctgg	attatgacgg	cactgttggt	cctgaaactt	60
ctatcagtaa	aatgccaggg	cctgaagtcc	tttctgtttt	gaacgctctc	tgtaatgac	120
caatgaacac	tgtattttatt	atcagtgggc	gggggagaaa	atcattaagt	gagtggcttt	180
cttcttgcaa	gaagcttgga	atagccgctg	agcatgggta	ttttataagg	tggaacagtg	240
cagccgagtg	ggaaaccagt	tcattgtctg	ctgatcttga	ttggaagaat	accgtggaac	300
ctataatgaa	ttcatcacaca	gaggcaactg	atggctcaag	catagagtac	aaggagagcg	360
ctttggtgtg	gcaccatcag	gatgcagacc	ctgatttttg	atcatgcca	gccaaggaat	420
tgttggctca	tctagagagt	gtgctcgcaa	atgaacctgc	agttgttaag	aggggacagc	480
aaattgtgga	ggtcaaacct	cagggagtaa	gcaaaggatt	ggttgcagag	a	531

&lt;210&gt; 189

&lt;211&gt; 329

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 189

atcctccaga	aatcagcttt	tggtgtctta	ctcctcaaaa	agtgaggcac	ttgccagttc	60
cttgagaaac	gagactacgt	cgtcagagga	tgtaagcatg	tagcgagcat	ttgtacgggt	120
ccgcccaca	gcgcagaga	aatagttttc	ccccttgaga	tgcagcacat	tccaagatat	180
tttatccgga	gatgacctcc	tcccactacc	actggcatgg	ttgttagttc	ttttctcagg	240
gttttgtaaa	ggtcggtgtg	tcttcaccgc	agacgctttt	gacccgcttt	tactagcttg	300
aagcttcaat	gaagatcggt	tctcccctg				329

&lt;210&gt; 190

&lt;211&gt; 503

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 190

acaaaaagcc	gagcgattct	tttggattat	gatggagcta	tgggggtcaac	aggatccaat	60
tccatcagtg	tgatacctac	tgctgagaca	gttgactca	ttaacagttt	gtgccgagat	120
cccaagaatg	ttgtcttcct	tgtcagtggg	aaggagcgag	ttatcctaag	taaattggttc	180
tcategtgtg	atagacttgg	gttagcagca	gagcatggct	atcttcttag	gccaaaccaa	240
gaaggagatt	gggaaacttg	tgtttcggtg	acagattttg	actggaaaca	gacagctgag	300
ccggttatga	gattatacat	ggaaacaact	gatggctcta	ctatagaaat	caaagagagt	360
tcaattgtgt	ggaactacca	gtgcgcagat	ccagattttg	gtttttgcca	ggcaaaggaa	420
cttttagatc	acctggaaa	gtttcttgca	aatgaacctg	tttctgtcaa	gagtggccaa	480
cacattgtgg	aagttaaacc	tca				503

&lt;210&gt; 191

&lt;211&gt; 398

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 191

acgacactag	aggatccaaa	gtctcgcaaa	aagggttgga	ggtcataaat	ccagaggatg	60
actatgtctg	gatacatgat	tatcattgga	gggggttgcc	tactttctta	agaaggaagt	120
tcataagctg	gaggatgggg	ttctttctcc	atagcccttt	tccatcatca	gagatatata	180
ggactcttcc	agnnaggag	gagatactca	aagcgcttct	taatgctgac	ctgattgggt	240
tccacaogtt	cgattatgct	cggcattttc	tatcctgttg	cagtagaatc	tgggngtg	300
agtaccagtc	gaaaaggggt	tatatcggt	tggaatatta	tggaagaaca	attggggtaa	360
agatcatgcc	tggtgggatc	cacatgggcc	agattcag			398

&lt;210&gt; 191

&lt;211&gt; 457

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 192

ggcaggctta	tgtctccgtc	aataagattt	tgcagatag	gatcatggaa	gtgattaatc	60
cagaggatga	tttcgtttgg	gtacacgatt	accatttgat	ggtggtgccg	actttcttga	120
ggaaaagggt	caatagagt	aagcttggt	tcttccttca	cagcccatc	ccctcttcag	180
agatttacia	gactctgccg	atcangagg	agcttctgan	ggctttcctg	aattctgatt	240
tgatagggtt	ccacactttc	gactatgcac	gccatttcct	gtcttggtgt	agtcggatgc	300
ttggtcttac	ctatgaatcg	aagaggggtt	atataggcct	ggactattat	ggtcggactg	360
ttagtatcaa	aattcttcca	gttgggatac	acatgggaca	attgcagtc	gttttaagcc	420
ttcccagac	tgaagccaag	gtggccgaac	taattaa			457

&lt;210&gt; 193

&lt;211&gt; 359

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 193

gcctgacctc	ggtggtcggt	tcaaccggtc	cttgtggcag	gcttatgtct	ccgtcaataa	60
gattttcgca	gataggatca	tggaagtgat	taatccagag	gatgatttcg	tttgggtaca	120
cgattaccat	ttgatgggtg	tgccgacttt	cttgaggaaa	aggttcaata	gagtgaagct	180
tggcttcttc	cttcacagcc	cattcccctc	ttcagagatt	tacaagactc	tgccgatcag	240
ggaggagctt	ctgagggcct	tcctgaattc	tgatttgata	gggttccaca	ctttcgacta	300
tgacgcctat	ttcctgtctt	gtttagtcg	gatgcttggt	cttacctatg	aatcgaaga	359

&lt;210&gt; 194

&lt;211&gt; 401

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 194

ttggcttctt	ccttcacagc	ccattcccct	cttcagagat	ttacaagact	ctgccgatca	60
gggaggagct	tctgagggct	ttcctgaatt	ctgatttgat	aggggtccac	actttcgact	120
atgcacgcca	tttctgtctt	tggttagtc	ggatgcttgg	tcttacctat	gaatcgaaga	180
ggggttatat	aggcctggac	tattatggtc	ggactgttag	tatcaaaatt	cttccagttg	240
ggatacacat	gggacaattg	cagtcggtt	taagccttcc	cgagactgaa	gccaagggtg	300
ccgaactaat	taagcagttt	ggtggtcggg	gtaggacaat	gttgctcggt	gtagatgaca	360
tggaatattt	taaaggaatt	agcttgaaac	tggtggccat	g		401

&lt;210&gt; 195

&lt;211&gt; 364

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 195

ccgactttct	tgaggaaaag	gttcaataga	gtgaagcttg	gcttcttcct	tcacagccca	60
ttcccctctt	cagagattta	caagactctg	ccgatcagg	aggagcttct	gagggctttc	120
ctgaattctg	atgtgatagg	gttccacact	ttcgactatg	cacgccattt	cctgtcttgt	180
tgtagtcgga	tgcttggtct	tacctatgaa	tcgaagagg	gttatatagg	cctggactat	240
tatggtcgga	ctggttagt	caaaattctt	ccagttggga	tacacatggg	acaattgcag	300
tcggttttaa	gccttcccga	gactgaagcc	aagggtggcg	aactaattaa	gcagtttggt	360
ggtc						364

&lt;210&gt; 196

&lt;211&gt; 400

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 196

gttttaaatgc	aacaatgact	gcacaagttg	atacacctgg	tagaagaggt	catgatcaaa	60
tcaaagagat	gaagctcctt	ttgcatcctg	atctgggtga	aaccctttca	gttttatgta	120
aggatccaaa	aactactata	gttgtgctta	gcggaagtga	aagaaatgtc	ttagatgaga	180
attttggcga	actcgacatg	tggttagcag	cagaaaatgg	tatgtttctg	cgccatacga	240
aaggagaatg	gatgatcaca	atgccagaat	atcttaatat	ggactggatg	gagagtgtaa	300
agcttgtttt	tgattatttc	aaagaaaggc	acctcgatcg	tatgtngagg	ttcgagaaac	360
ctcttttagtt	tggacctata	agtatgcaga	tgttgaattt			400

&lt;210&gt; 197

&lt;211&gt; 298

&lt;212&gt; DNA

<213> *Eucalyptus grandis*

&lt;400&gt; 197

taaaaccgnt	agcagcaagt	gctgtccatt	acaatgggtg	attggagtat	gatgtccaca	60
gtttgtatgg	tttctcgcaa	tcaattgcta	ctcacaaagc	actccaaggg	ctccagggaa	120
agagaccctt	catattgtcg	cgctcgacgt	acgtgggctc	cggaagtat	gtngctcact	180
ggaccggaga	taaccaaggn	aattgggaaa	atctgaagta	ttccatctcc	actatgctga	240
atttcggcat	attcggagtg	ccgatggctg	gtgcagacat	atgcgggttc	taccccaa	298

&lt;210&gt; 198

&lt;211&gt; 402

&lt;212&gt; DNA

<213> *Eucalyptus grandis*

&lt;400&gt; 198

ctggaccgga	gataaccaag	gcaattggga	aaatttgaag	tattccatct	ccactatgct	60
gaatttttggc	atattcggag	tgccgatggg	cggtgcagac	atatcgggct	tctaccggc	120
cccgaactgag	gagctttgca	accgctggat	tgaaagtcgg	tgcttntac	ccntntttt	180
tngggggggn	ttncctnntt	tttttttnc	cnaaaagggg	tttttaatgg	gactcagtag	240
ccgaatctgc	taggaatgct	cttggcatga	gatataggct	cctaccttac	ttgtacactc	300
tcaattacca	agctcatacg	acgggagccc	cgattgcacg	gcctcttttc	ttttcattcc	360
ccgattacgt	ggagagttac	ggattgagca	cccagttttt	gc		402

&lt;210&gt; 199

&lt;211&gt; 441

&lt;212&gt; DNA

<213> *Pinus radiata*

&lt;400&gt; 199

atcgacccag	ggattggcgt	taacacgagc	tacgggacgt	tccagcgagg	aatggcggac	60
gacgttttca	taaagcacga	cgggagtcg	ttcttgggtc	aggtgtggcc	cggcgccgtg	120
tactttccgg	acttctgaa	cccaaagacg	gtggatttct	gggcccagca	gatctcccgc	180
ttccacaaa	tgggtcccgt	ggacggtctc	tggatcgaca	tgaacgaggt	ctccaatttc	240
tgcagtggca	agtgtcccat	ccccacaaac	cggagctgcc	cgggcacggg	tttcccatgg	300
gactgctgcc	tgcactgcac	aaacatcacc	gccacccgat	gggacgtgcc	accctaccag	360
atcaacgcct	cgggtaccca	ggtcccgtcg	gggttcaaga	ccatcgccac	cagctccgtc	420
cactacaacg	gcgtcctcga	g				441

&lt;210&gt; 200

&lt;211&gt; 481

&lt;212&gt; DNA

<213> *Pinus radiata*

&lt;400&gt; 200

ctataacaaa	tctcagatag	ctctggatgt	tatatggaac	gatgatgacc	acatggatgg	60
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agccaaagac ttcacccttg accctatcaa ctatcctgaa tataagctgc gtcccttcct 120
tgaccgaatt catgccaatg gaatgagata tgctcgtcct atcgacccag ggattggcgt 180
taacacgagc tacgggacgt tccagcgagg aatggcggac gacgttttca taaagcacga 240
cgggagtcgg ttcttgggtc aggtgtggcc cggcgccgtg tactttccgg acttcctgaa 300
cccaaagacg gtggatttct gggccgacga gatctcccgc tccaccaaaa tgggtccccgt 360
ggacgggtctc tggatcgaca tgaacgaggt ctccaatttc tgcagtggca agtgctccat 420
ccccacaaac cggagctgcc cgggcacggt ttcccatggg actgctgcct cgactgcaca 480
a 481

```

&lt;210&gt; 201

&lt;211&gt; 484

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 201

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ggtggtgctg tgttcaaagt agactggatg agctccgtac aagttggtac caggagcaac 60
accataagca tcacggttcc aaatagtcct agtagtgttg gaaggatcga ggcggaaagt 120
gttggtatgt tcgccgagac catagatgtt cgcgttatta gggacgacgg tcttaagtgc 180
gagatattgc tcctcaaaca caagcgtgct acccttggtg tcgaacaaca cctcgccaga 240
ctttttacgg acaacgctga acgagaacgg gctcggaaca tatttgaaga cgatatctgc 300
cgaggacgcc gtgaccttgg atgttgacg cgggaaaacc tcctcgggga cttcatatcg 360
cttgccccgc gaatcaccaa tcttgacatg aactcgagac tcgtcttcgt acgtaacctg 420
caacttgagc tgctgaacgt cactgccgta gatccacag gtagcggcga gagtcaggtc 480
agcg 484

```

&lt;210&gt; 202

&lt;211&gt; 418

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 202

```

caatgcaggt ggggatacaa gaacgtatca gacataacga atgtggtgga aaactataac 60
aaatctcaga tacctctgga tgttatatgg aacgatgatg accacatgga tggagccaaa 120
gacttcaccc ttgaccctat caactatcct gaatataagc tgcgtccctt ccttgaccga 180
attcatgcca atggaatgag atatgtcgtc cttatcgacc cagggtattg cgtaaacacg 240
agctacggga cgttccagcg aggaatggcg gacgacgttt tcataaagca cgacgggagt 300
ccgttcttgg gtcaggtgtg gcccgccgcc gtgtactttc cggacttcct gaacccaaag 360
acggtggatt tctgggccga cgagatctcc cgcttccacc aaatgggtccc cgtggacg 418

```

&lt;210&gt; 203

&lt;211&gt; 395

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 203

```

caatgcaggt ggggatacaa gaacgtatca nacataacga atgtggtgga aaactataac 60
aaatctcaga tacctctgga tgttatatgg aacgatgatg accacatgga tggagccaaa 120
gacttcaccc ttgaccctat caactatcct gaatataagc tgcgtccctt ccttgaccga 180
attcatgcca atggaatgag atatgtcgtc cttatcgacc cagggtattg cgtaaacacg 240
agctacggga cgttccagcg aggaatggcg gacgacgttt tcataaagca cgacgggagt 300
ccgttcttgg gtcaggtgtg gcccgccgcc gtgtactttc cggacttcct gaacccaaag 360
acggtggatt tctgggccga cgagatctcc cgctt 395

```

&lt;210&gt; 204

&lt;211&gt; 390

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

## &lt;400&gt; 204

cgtccttatac	gacccagggga	ttggcggttaa	cacgagctac	ggcacgttcc	agcgaggaat	60
ggcggacgac	gttttcataa	agcacgacgg	gagtcggttc	ttgggtcagg	tgtggcccgg	120
cgccgtgtac	tttccggact	tcctgaaccc	aaagacggtg	gatttctggg	ccgacgagat	180
ctcccgttcc	caccaaagtg	tccccgtgga	cggtctctgg	atcgacatga	acgaggtctc	240
caatttctgc	agtggcaagt	gctccatccc	cacaaaccgg	agctgcccgg	gcacgggttt	300
cccatgggac	tgctgcctcg	actgcacaaa	catcacccgc	acccgatggg	acgtgccacc	360
ctaccagatc	aacgcctcgg	taccaagtc				390

## &lt;210&gt; 205

## &lt;211&gt; 245

## &lt;212&gt; DNA

## &lt;213&gt; Eucalyptus grandis

## &lt;400&gt; 205

ccaccgcaga	atccccccgg	cagtcctcgg	aatcatgttc	ttgtctgggtg	gacaatctga	60
agttgaagcc	accctgaact	tgaatgccat	gaaccaatcg	ncaaaccctg	ggcacgnatc	120
tttctcatat	gcccagagccc	tccagaacac	ctgcttgaag	acatggggag	gaagaccoga	180
gaacgtgaag	ccagcccagg	aaaccttgct	tgtccgcgcc	aagggcaaat	ctcttgctca	240
aactt						245

## &lt;210&gt; 206

## &lt;211&gt; 510

## &lt;212&gt; DNA

## &lt;213&gt; Eucalyptus grandis

## &lt;400&gt; 206

tgagaagcga	gagaagcaaa	gcaagaagca	atggcctctg	cttctgcaac	gctgctcaag	60
tcattccccag	tccttgacaa	gtccgagtg	gtgaagggtc	agtctatccg	ccacgcctcg	120
gccaccaccg	tgcgtgcca	tcctgccacc	gcctctgtcc	tcactgtcaa	agccagtaat	180
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atgggtgatg	tacttggtga	gcagaacatc	gtccccggta	tcaaagtcga	caaggggttg	480
tgctttggcg	gtttaacaat	gagtctgggtg				510

## &lt;210&gt; 207

## &lt;211&gt; 413

## &lt;212&gt; DNA

## &lt;213&gt; Eucalyptus grandis

## &lt;400&gt; 207

tggtgccttt	ggctgggtcc	aacgacgagt	cctggtgcca	agggctcgat	ggattagcat	60
cccgcacggc	tgcttactac	cagcaggggtg	cacgctttgc	taaatggaga	actgtggtga	120
gcatccccaa	tgccccatct	gccctggccg	tgaagggaagc	cgccctgggtg	cttggccgct	180
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tctactacat	ggctgagaac	aatgtactgt	tcgagggcat	cctcttgaag	cccagcatgg	360
tcactcccg	cgctgagtgc	aaagagaggg	ccaactccca	gcaagtggcc	gag	413

## &lt;210&gt; 208

## &lt;211&gt; 434

## &lt;212&gt; DNA

## &lt;213&gt; Eucalyptus grandis

&lt;400&gt; 208

gctgtccctg	ccgttgtgtt	tttgtctggt	gggcagagtg	aggaggaggc	aaccctcaac	60
ctcaatgcca	tgaacaagct	caagggcaag	aaaccatggt	ctctttcctt	ctcctttgga	120
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tactaagggtg	cttttgcctg	cggtattttc	tgcttttcat	ttgagaaaaa	taggtgtctg	360
gagatattcct	atttttgtac	taggataata	agcatgactg	tactgtactg	gtaatgcatt	420
ttcattgatg	ttgt					434

&lt;210&gt; 209

&lt;211&gt; 350

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 209

actttagtc	cttaacatgg	agagactcgg	aagcaccctc	gccaaagtga	gcatcaccct	60
tgtatgttcc	aagagttgcg	tccgagttgg	ccttgcatct	tctgagaagt	gcagcctgtg	120
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caggaggcac	tgttcgctgc	agggcacgaa	cggtgtactc	agcaatgacc		350

&lt;210&gt; 210

&lt;211&gt; 455

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 210

gcgaaaaaca	atggcctcgg	cttctctcct	caagtcatct	cccgtgctcg	acaagtcgga	60
gtttgtgaag	ggcacgcaga	ctctccgcac	cccattctct	gccgcgctcc	gctaccaccc	120
caccaccgcc	ccctccgctc	tcgtcgtcaa	agccagtgcc	tatgctgatg	agctcgtcaa	180
gactgcgaaa	acagttgcat	cacccgaggag	aggaatcctg	gccatggacg	agtcaaacgc	240
aacctgcggg	aagcgtttgg	cgtcgatcgg	gctagagaa	accgaggcca	accgccaggc	300
ctacaggaca	ctcctgggca	gtgctccggg	gcttggccag	tacatctctg	gtgctatcct	360
cttcgaggag	actctctacc	aatccaccac	cgacggccgc	aagatggtcg	acgtcctcgt	420
tgagcagaac	attgtccctg	gtattaaagt	cgaca			455

&lt;210&gt; 211

&lt;211&gt; 509

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 211

ggagggagaa	gctttgagaa	gagataaggt	aacacagttt	ttgcttgccg	ggctgtgctc	60
gagtttggcc	tttgcaagcg	aaaaacaatg	gcctcggtct	ctctcctcaa	gtcatctccc	120
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gccgtccgct	accaccccac	caccgcccc	tccgtctctg	tcgtcaaagc	cagtgcctat	240
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gaggccaacc	gccaggccta	caggacactc	ctggtcagtg	ctccggggct	tggccagtac	420
atctctgggtg	ctatcctctt	cgaggagact	ctctaccaat	ccaccaccga	cggccgcaag	480
atgggtcgacg	tcctcggtga	gcagaacat				509

&lt;210&gt; 212

&lt;211&gt; 364

&lt;212&gt; DNA

## &lt;213&gt; Eucalyptus grandis

&lt;400&gt; 212

atttgaagtt gctcagaagg tttgggctga ggttttctac tacctggctg agaacaatgt	60
catgtttgag ggtatcctcc ttaagcccag catggctact cctgggtgccg agtgcaagga	120
caaggccact cctcaacaag tcgctgaata caccctcaag cttctccacc gcagaatccc	180
cccggcagtc cctggaatca tgttcttgtc tgggtggacaa tctgaagttg aagccaccct	240
gaacttgaat gccatgaacc aatcgccaaa cccgtggcac gtatctttct catacgcccg	300
agccctccag aacacctgct tgaagacatg gggaggaaga cccgagaacg tgaagccagc	360
ccag	364

&lt;210&gt; 213

&lt;211&gt; 372

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 213

ctgaagttga agccaccctg aacttgaatg ccatgaacca atcgccaaac ccggtggcacg	60
tatctttctc atacgccga gccctccaga acacctgctt gaagacatgg ggaggaagac	120
ccgagaacgt gaagccagcc caggaaacct tgcttgcctg cgccaaggcc aactctcttg	180
ctcagcttgg caagtacaca ggtgaggcg agtccgagga ggccaagaaa ggaatgttcg	240
tcaagggcta cgtgtattaa gctgttctact gtaggtggaa gtggatgatc aaagttggga	300
gacttaagaa ttgatccctc tcagcgtgtt atgattatag ccacggagac tatttttgca	360
catcgaatgt ac	372

&lt;210&gt; 214

&lt;211&gt; 471

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 214

atattaacct cctcatataa aaatataaaa cttcataaag aagtcgcaaa cattcaacca	60
tcacaacatc aatgaaaatg cattaccagt acagtcatgc ttattatcct agtacaaaaa	120
taggatatct ccagacacct atttttctta aaaggaaaag cagaaaatac cgccagcaaa	180
aaagcacctt agtacttgta atccttaaca tggagacttt cagcagctcc ctgcgcaagc	240
tttgcgtcac ccttgtaagt accaagagtt gcctctgagt ttgccttgca cctagttaat	300
aatgcagcct gtgccttggg cacgttctcc tcctttccag cccaagactt caaagtgtct	360
tgctgaagag cccgtccaaa ggagaaggaa agagaccatg gtttcttgcc cttgagcttg	420
ttcatggcat tgagggtgag ggttgctcc tcctcactct gccaccaga c	471

&lt;210&gt; 215

&lt;211&gt; 465

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 215

acggtcagta atccacgaaa caaacgaagc acaagtaata gtatcgcagc aaagcctaga	60
tccgcccctc agtacttgta gtccttaaca tggagagact cggaagcacc ctgcgcaagt	120
tgagcatcac ccttgatgtg tccaagagtt gcgtccgagt tggccttgca tcttctgaga	180
agtgcagcct gtgccttggg aatgttctct tcctttccag cccacgcctt caaggtgctc	240
tgctgtaggg cccgacaaa agagaaggag aggtcccatg gcttcttgcc cttgagcttg	300
ttcatggcat tgagggtcaa agtggcctcc tcctcactct gccaccgga caagaagaca	360
atggccggaa cagcaggagg cactgttcgc tgcagggcac gaacggtgta ctcagcaatg	420
acctccggag caaccttggg ggcattctgat ccgggggtga ccatg	465

&lt;210&gt; 216

&lt;211&gt; 484

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 216

ctcatataaa	aatataaaa	ttcataaaga	agtcgcaaac	attcaacccat	cacaacatca	60
atgaaaatgc	attaccagta	cagtcatgct	tattatccta	gtacaaaaat	aggatatctc	120
cagacaccta	tttttctcaa	atgaaaagca	gaaaataccg	ccagcaaaaa	agcaccttag	180
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gccttgggca	cgttctctc	ctttccagcc	caagacttca	aagtgtctctg	ctgaagagcc	360
cgtccaaagg	agaaggaaa	agaccatggt	ttcttgccct	tgagcttggt	catggcattg	420
aggttgaggg	ttgcctctc	ctcactctgc	ccaccagaca	aaaacacaac	ggcagggaca	480
gccg						484

&lt;210&gt; 217

&lt;211&gt; 362

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 217

cgccctcag	tacttgtagt	ccttaacatg	gagagactcg	gaagcaccct	cgccaagttg	60
agcatcacc	ttgtatgttc	caagagttgc	gtccgagttg	gccttgcatc	ttctgagaag	120
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catggcattg	aggttcaaag	tggcctctc	ctcactctgc	ccaccggaca	agaagacaat	300
ggccggaaca	gcaggaggca	ctgttcgctg	cagggcacga	acggtgtact	cagcaatgac	360
ct						362

&lt;210&gt; 218

&lt;211&gt; 395

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 218

ccttaacatg	gagagactcg	gaagcaccct	cgccaagttg	agcatcacc	ttgtatgttc	60
caagagttgc	gtccgagttg	gccttgcatc	ttctgagaag	tgcagcctgt	gccttgggaa	120
tgttctcttc	ctttccagcc	cacgccttca	aggtgctctg	ctgtagggcc	cgaccaaag	180
agaaggagag	gtcccatggc	ttcttgccct	tgagcttggt	catggcattg	aggttcaaag	240
tggcctctc	ctcactctgc	ccaccggaca	agaagacaat	ggccggaaca	gcaggaggca	300
ctgttcgctg	cagggcacga	acggtgtact	cagcaatgac	ctccggagca	accttggggg	360
cattgatccg	ggggtgacca	tgttgggctt	caaca			395

&lt;210&gt; 219

&lt;211&gt; 416

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 219

cgccctcag	tacttgtagt	ccttaacatg	gagagactcg	gaagcaccct	cgccaagttg	60
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tgcagcctgt	gccttgggaa	tgttctcttc	ctttccagcc	cacgccttca	aggtgctctg	180
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catggcattg	aggttcaaag	tggcctctc	ctcactctgc	ccaccggaca	agaagacaat	300
ggccggaaca	gcaggaggca	ctgttcgctg	cagggcacga	acggtgtact	cagcaatgac	360
ctccggagca	accttggggg	catctgatcc	gggggtgacc	atgttgggct	tcaaca	416

&lt;210&gt; 220

<211> 452  
 <212> DNA  
 <213> Eucalyptus grandis

<400> 220  
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 tgagcatcac ccttgatgtg tccaagagtt gcgtccgagt tggccttgca tcttctgaga 180  
 agtgcagcct gtgccttggg aatgtttctt tcctttccag cccacgcctt caaggtgctc 240  
 tgctgtaggg cccgaccaa agagaaggag aggtcccatg gcttcttgcc cttgagcttg 300  
 ttcattggcat tgaggttcaa agtggcctcc tctcactct gccaccgga caagaagaca 360  
 atggccggaa cagcaggagg cactgttcgc tgcagggcac gaacggtgta ctcagcaatg 420  
 acctccggag caaccttggg ggcattctgat cc 452

<210> 221  
 <211> 289  
 <212> DNA  
 <213> Eucalyptus grandis

<400> 221  
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 cacccttgna tggtccaaga gttgcgtccg agttggcctt gcatcttctg agaagtgcag 120  
 cctgtgcctt gggaatgttc tcttcctttc cagcccacgc cttcaagggtg ctctgctgta 180  
 gggcccgacc aaaagagaag gagaggctcc atggcttctt gcccttgagc ttgttcatgg 240  
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<210> 222  
 <211> 460  
 <212> DNA  
 <213> Eucalyptus grandis

<400> 222  
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 ccaaattccat tgcattctct ggtcgttgta tccttgccat tgatgagtca aatgcaacat 180  
 gtgggaaaaa gttagcatcc atcgggttgg acaacactga ggtcaatcgt caagcttata 240  
 gacaacttct gttgaccacg cctgggtctg gtgaatacat ctctgggtgc attttgtttg 300  
 aggagacact ttaccaatcg acaacagatg ggaagaaatt tgttgactgc ctgcgtgagg 360  
 agaaaattgt tccaggcatt aaagtgtgaca aggggttgggt tcctcttctt ggatccaata 420  
 atgaatcctg gtgccaaggc ttggatggat tggcttcaag 460

<210> 223  
 <211> 373  
 <212> DNA  
 <213> Eucalyptus grandis

<400> 223  
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 tgcagcctgt gccttgggaa tgttctcttc cttccagcc cagccttca aggtgctctg 180  
 ctgtagggcc cgaccaaag agaaggagag gctccatggc ttcttgccct tgagcttggt 240  
 catggcattg aggttcaaag tggcctctc ctcactctgc ccaccggaca agaagacaat 300  
 ggccggaaaca gcaggaggca ctgttcgctg cagggcacga acggtgtact cagcaatgac 360  
 ctccggacaa cct 373

<210> 224  
 <211> 524

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 224

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cccgcgcgt	ctccgccccg	atccgcgcgc	ccgcttactc	cgacgagetc	gtccagaccg	120
ccaaatccat	tgcattctct	ggtcgtggta	tccttgccat	tgatgagtca	aatgcaacat	180
gtgggaaaag	gttagcatcc	atcgggttgg	acaacactga	ggtaaatcgt	caagcttata	240
gacaacttct	gttgaccacg	cctgggtctg	gtgaatacat	ctctgggtgc	attttgtttg	300
aggagacact	ttaccaatcg	acaacagatg	ggaagaaatt	tggtgactgc	ctgcgtgagg	360
agaaaattgt	tccaggcatt	aaagttgaca	agggtttggg	tcctcttctc	ggatccaata	420
atgaatcctg	gtgccaaggc	ttggatggat	tggcttcaag	gtccgctgaa	tactacaagc	480
aagggtgctg	ttttgccaaa	tggaggacag	tggttagcat	tcct		524

&lt;210&gt; 225

&lt;211&gt; 332

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 225

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tccgcccctc	agtacttgta	gtccttaaca	tggagagact	cggaagcacc	ctcgccaagt	120
tgagcatcac	ccttgatgt	tccaagagtt	gcgtccgagt	tggccttgca	tcttctgaga	180
agtgcagcct	gtgccttggg	aatgttctct	tcctttccag	cccacgcctt	caaggtgctc	240
tgctgtaggg	cccgacaaa	agagaaggag	aggctccatg	gcttcttgcc	cttgagcttg	300
ttcatggcat	tgaggttcaa	agtggcctcc	tc			332

&lt;210&gt; 226

&lt;211&gt; 362

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 226

cgtccctcag	tactttagt	ccttaacatg	gagagactcg	gaagcaccct	cgccaagttg	60
agcatcacc	ttgtatgtc	caagagttgc	gtccgagttg	gccttgcatc	ttctgagaag	120
tgagcctgt	gccttgggaa	tggtctcttc	ctttccagcc	cacgccttca	aggtgctctg	180
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catggcattg	aggttcaaag	tggcctcttc	ctcactctgc	ccaccggaca	agaagacaat	300
ggccggaaca	gcaggaggca	ctgttcgctg	cagggcacga	acgggtgtact	cagcaatgac	360
ct						362

&lt;210&gt; 227

&lt;211&gt; 506

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 227

gtcactcccc	gcgctgagt	caaagagagg	gccactcccc	agcaagtgcg	cgagtacacc	60
ctcaagctcc	tccaccgcag	gatcccgcct	gccgttcccc	gaatcatggt	cttgtctggt	120
gggcaatccg	aggtcgaagc	aacctgaac	ctgaacgcga	tgaaccagtc	cccgaaccca	180
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atgttcgtca	agggatacgt	gtactaaggc	gatgcactga	aactccatga	gctcagaaga	420
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agtagaacag	ctaattcctc	cttctc				506

<210> 228  
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 <212> DNA  
 <213> Eucalyptus grandis

<400> 228  
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 atggattggc ttcaagggtcc gctgaataact acaagcaagg tgctcgtttt gccaaatgga 120  
 ggacagtggg tagcattcct tgtgggtccct ctgctcttgc agtaaaagaa gctgctggg 180  
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<210> 229  
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 <213> Eucalyptus grandis

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 gggcaatccg aggtcgaagc aaccctgaac ctgaacgcga tgaaccagtc cccgaaccca 180  
 tggcacgtgt ccttctccta cgctagagcc ctccagaaca cctgcttgaa gacgtgggga 240  
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 tccctcgccc agctcggcaa gtacaccggg gaaggcgagt ctgaggaggc caagaaggga 360  
 atgttcgtca agggatacgt gtactaaggc gatgcactga aactccatga gctcagaaga 420  
 tgatcacagg gtttagttat gataatgatg 450

<210> 230  
 <211> 417  
 <212> DNA  
 <213> Eucalyptus grandis

<400> 230  
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 gggcaatccg aggtcgaagc aaccctgaac ctgaacgcga tgaaccagtc cccgaaccca 180  
 tggcacgtgt ccttctccta cgctagagcc ctccagaaca cctgcttgaa gacgtgggga 240  
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 tccctcgccc agctcggcaa gtacaccggg gaaggcgagt ctgaggaggc caagaaggga 360  
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<210> 231  
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 <212> DNA  
 <213> Eucalyptus grandis

<400> 231  
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 ccgccccgat ccgcgcggcc gcttactccg acgagctcgt ccagaccgcc aaatccattg 180  
 catctcctgg tcgtggtatc cttgccattg atgagtcaaa tgcaacatgt gggaaaagg 240  
 tagcatccat cgggttgagc aatactgagg tcaatcgta agcttataga caacttctgt 300  
 tgaccacgcc tggctctggg gaatacatct ctggtgccat tttgtttgag gagacacttt 360  
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aagctgcgtg gggacttgca cgatatgctg gcattctctca ggataatggc cttgtgccca 660  
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<210> 232  
 <211> 435  
 <212> DNA  
 <213> Eucalyptus grandis

<400> 232  
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 aacagctccc tcgccaagct ttgcgtcacc cttgtaagta ccaagagttg cctctgagtt 180  
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 aacagtgtac tcagc 435

<210> 233  
 <211> 352  
 <212> DNA  
 <213> Eucalyptus grandis

<400> 233  
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<210> 234  
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 <212> DNA  
 <213> Pinus radiata

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 gtgctgagca caaggagaga gcaacccccg 330

<210> 235  
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 <212> DNA  
 <213> Pinus radiata

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 ggttgaggca acattgaatt tgaatgcaat gaaccaaagc ccaaaccat ggcatgtttc 180  
 cttttcatat gcacgtgctt tgcagaatac atctctcaag acctgaaagg gtcttccaga 240  
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 g 301

<210> 236  
 <211> 368  
 <212> DNA  
 <213> Pinus radiata

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 acaaggagag agcaaccccc gaaaagggtg cagagtacac tctaaaaatg cttagagga 360  
 gggtgcc 368

<210> 237  
 <211> 423  
 <212> DNA  
 <213> Pinus radiata

<400> 237  
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 cgggtctcgg acagtacatc tccggcgcca ttctcttca ggaactctg taccagtcca 120  
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 gcgtacagtt gtgagcattc ccaacggccc ctctgctctg gccgtgaaag aagccgcatg 360  
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<210> 238  
 <211> 352  
 <212> DNA  
 <213> Pinus radiata

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<210> 239  
 <211> 427  
 <212> DNA  
 <213> Pinus radiata

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 tcccggggaga ggcattcctg cgatggacga gtccaacgcc acctgtggga aacggctggc 120  
 gtccatcgagg cttgagaaca cggaggcgaa ccgacaggca tacaggcagc tgctcgtcag 180  
 cgcgcgggggt ctccgacagt acatctccgg cgccattctc ttcgaggaaa ctctgtacca 240  
 gtccagcacc gaaggcaaga agattgtaga catcctcgtg caacagaaca tagtccccgg 300  
 catcaaagtt gacaagggtc tggttccttt ggctggttca aacgacgaat cttgggtgcca 360  
 aggctagac ggcctcgcac caaggtgcgc tgagtattat aagcaaggag ctcgcttcgc 420  
 caaatgg 427

<210> 240  
 <211> 470  
 <212> DNA  
 <213> Pinus radiata

<400> 240  
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 ggcaacattg aatttgaatg caatgaacca aagcccaaac ccatggcatg tttccttttc 180  
 atatgcacgt gccttgcaga atacatctct caagacctgg aagggtcttc cagagaatgt 240  
 tgaagcagct cagagggcgc ttcttattcg ggccaaggct aattctctgg ccagcttgg 300  
 gcgatactct gctgaagggt aaagtgagga gtctaagaag ggaatgttcg ttaagggata 360  
 cacatattaa gaatgtgggt catagttttc ttacgggaag aactcgttca atgcggatag 420  
 gttaagcttt tatgtttatt tanttggcac ttacaatcct gaacttttta 470

<210> 241  
 <211> 396  
 <212> DNA  
 <213> Pinus radiata

<400> 241  
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 aaatttggtg actgtttgcg agagcagaat attatgcccg gcatcaaagt tgacaagggg 120  
 ttagttccac tgccaggatc aaacaatgaa tcttggtgcc aggggttggg tggattagcc 180  
 tcaagatctg ctgagtacta caaacagggt gcaagatttg ctaaatggcg aacagttgtc 240  
 agcataccca atgggccttc ggagtttagct gtgaaggaag ctgcgtgggg acttgacagt 300  
 tatgccgcta tctctcagga caatggtctt gtgcccattg tggagccaga gattcttctg 360  
 gatggagacc attgcattga cagaagcttg aagtgg 396

<210> 242  
 <211> 273  
 <212> DNA  
 <213> Pinus radiata

<400> 242  
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 cacaaggaga gagcaacccc cgaaaaggtt gcagagtaca ctctaaaaat gcttaagagg 120  
 aggggtgccac cagctgttcc tgggggttatg ttcttgtctg gaggacagtc tgaggttgag 180  
 gcaacattga atttgaatgn aatgaaccaa agcccaaacc catggcatgt ttccttttca 240  
 tatgcacgtg ccttgcagaa tacatctctc aag 273

<210> 243  
 <211> 557  
 <212> DNA  
 <213> Pinus radiata

<400> 243  
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 cattgtggag ccagagattc ttctggatgg agaccattgc attgacagaa gccttgaagt 120  
 ggcggagaaa gtctgggctg aggttttctt ttacttggca cagaacaatg tgtgtttga 180  
 gggatatttg ttaaagccca gtatggtgac tcctggtgct gagcacaagg agagagcaac 240  
 ccctgaaaag gttgcagagt acactctaaa aatgcttaag aggaggggtgc caccagctgt 300  
 tcctgggggtt atgttcttgt ctggaggaca gtctgaggtt gaggcaactt tgaatttgaa 360  
 tgcaatgaac caaagcccaa atccatggca tgtttccttt tcatatgcac gtgccttgca 420  
 gaatacatct ctcaagacct ggaaggggtct tccagagaat gttgaagcag ctgagagggc 480  
 gcttcttatt cgggccaagg ctaattctct ggcccagctt gggcgatact ctgctgaagg 540  
 tgaaagttag gagtcta 557

<210> 244  
 <211> 593  
 <212> DNA  
 <213> Pinus radiata

<400> 244  
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 ccagcaccga aggcaagaag attgtagaca tcctcgtgca acagAACATA gtccccggta 120  
 tcaaagttga caaggggtctg gttccttttg ctggttcaaa cgacgaatct tgggtgccaa 180  
 gcctagacgg cctcgcatca aggtgcgctg agtattataa gcaaggagct cgcttcgcca 240  
 aatggcgtag agttgtgagc attcccaacg gccctctgc tctggccgtg aaagaagccg 300  
 catgggggtct cgcccgtac gcggcaattg ctcaggacaa cggctctggt cccatagtgg 360  
 agccagagat catgttggat ggagaacacg gcattgagag gactttcgaa gtagcgctga 420  
 aggtgtggtc cgaggtgttc ttctacctag cagagaacaa cgtgctgttc gaaggcattc 480  
 tgctgaagcc aagcatgggt acccctgggt ccggaatgta aggagagggc cagtcccgaa 540  
 actgttgccc aatataccct gaaccttctc cgaagaagaa ttccaccggc cgt 593

<210> 245  
 <211> 485  
 <212> DNA  
 <213> Pinus radiata

<400> 245  
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 ccagcaccga aggcaagaag attgtagaca tcctcgtgca acagAACATA gtccccggta 120  
 tcaaagttga caaggggtctg gttccttttg ctggttcaaa cgacgaatct tgggtgccaa 180  
 gcctagacgg cctcgcatca aggtgcgctg agtattataa gcaaggagct cgcttcgcca 240  
 aatggcgtag agttgtgagc attcccaacg gccctctgc tctggccgtg aaagaagccg 300  
 catgggggtct cgcccgtac gcggcaattg ctcaggacaa cggctctggt cccatagtgg 360  
 agccagagat catgttggat ggagaacacg gcattgagag gactttcgaa gtagcgctga 420  
 aggtgtggtc cgaggtgttc ttctacctag cagagaacaa cgtgctgttc gaaggcattc 480  
 tgctg 485

<210> 246  
 <211> 477  
 <212> DNA  
 <213> Pinus radiata

<400> 246  
 caggatcaaa caatgaatct tgggtgccagg gtttggatgg attagcctca agatctgctg 60  
 agtactacaa acaggggtgca agatttgcta aatggcgaa agttgtcagc ataccCAATG 120  
 ggccttcgga gttagctgtc aaggaagctg cgtggggact tgcacgttat gctgctatct 180  
 ctcaggacaa tgggtcttggt cccattgtgg agccagagat tcttctggat ggagaccatt 240  
 gcattgacag aagccttgaa gtggcggaga aagtctgggc tgaggttttc ttttacttgg 300  
 cacagaacaa tgtgttgttt gagggatatt tggttaaagcc cagtatgggtg actcctgggtg 360  
 ctgagcacia ggagagagca acccctgaaa aggttgca ga gtacactcta aaaatgctta 420  
 agaggagggt gccaccagct gtcttgggggt tatgttcttg tctggaggac agtctga 477

<210> 247  
 <211> 337  
 <212> DNA  
 <213> Pinus radiata

<400> 247  
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 cgaatcttgg tgccaaggcc tagacggcct cgcacaaagg tgcgctgagt attataagca 120

aggagctcgc	ttcgccaaat	ggcgtacagt	tgtgagcatt	cccaacggcc	cctctgctct	180
ggccgtgaaa	gaagccgcat	ggggctctgc	ccgctacgcy	gcaattgctc	aggacaacgg	240
tctggttccc	atagtggagc	cagagatcat	ggtggatgga	gaacacggca	ttgagaggac	300
tttcgaagta	gcgctgaagg	tgtggtccga	ggtgttc			337

&lt;210&gt; 248

&lt;211&gt; 452

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 248

gttttctttt	acttggcaca	gaacaatgtg	ttgtttgagg	gtattttgtt	aaagccaagt	60
atggtgactc	ctggtgctga	gcacaaggag	agagcaaccc	ccgaaaaggt	tgcagagtac	120
actctaaaaa	tgcttaagag	gagggtgcca	ccagctgttc	ctggggttat	gttcttgtct	180
ggaggacagt	ctgaggttga	ggcaacattg	atttgaatgc	aatgaaccaa	agcccaaacc	240
catggcatgt	ttccttttca	tatgcacgtg	ccttgacaga	tacatctctc	aagacctgga	300
agggtcttcc	agagaatgtt	gaagcagctc	agagggcgct	tcttattcgg	gccaaggcta	360
attctctggc	ccagcttggg	cgataactctg	cttgaagggtg	aaagtgagga	gtctaagaag	420
ggaatgttcg	ttaagggata	cacatattaa	ga			452

&lt;210&gt; 249

&lt;211&gt; 358

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 249

cagaatacac	agtaagggct	cttcagagga	ctgtgccacc	tgcagtggcc	ggcataatgt	60
ttctatctgg	tgggcagagt	gaggaggagg	ccaccttgaa	cttgaatgcc	atgaacaagc	120
tgcagaccaa	gaagccctgg	acattgnat	tctcctttgg	ccgggctctt	caggccagca	180
ctttgaagac	atgggctgga	aaggatgaga	atattcctgc	ggctcaggct	gccttgttat	240
ctcgatgcaa	ggccaattct	gatgccactt	tgggcaagta	tgcagggtgat	tctgctaagg	300
gcaatgggtg	ttctgagagc	cttcatgtca	aggactataa	gtattgattg	atgaccac	358

&lt;210&gt; 250

&lt;211&gt; 341

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 250

aaaactatga	cccacattct	taatattgtg	atcccttaac	gaacattccc	ttcttagact	60
cctcactttc	accttcagca	gagtatcgcc	caagctgggc	cagagaatta	gccttggccc	120
gaataagaag	cgccctctga	gctgcttcaa	cattctctgg	aagacccttc	caggctctga	180
gagatgtatt	ctgcaaggca	cgtgcatatg	aaaaggaaaac	atgccatggg	tttgggcttt	240
ggttcattgc	attcaaattc	aatgttgctt	caacctcaga	ctgtcctcca	gacaagaaca	300
taaccccagg	aacagctggg	ggcaccctcc	tcttaagcat	t		341

&lt;210&gt; 251

&lt;211&gt; 408

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 251

gaaattccag	agaaagacct	tgtgatctat	gaaatgagtg	ttcgatcctt	cacagcagac	60
aaatcaagtg	ggttggaacc	cagtatacgt	ggaagctatc	ttgggtgttat	tgaaaagatt	120
cctcatcttc	tagaacttgg	cattaatgca	gtggaattat	taccagtgtt	tgagtttgac	180
gagtttgaat	ttcaaaggca	tccaaatcct	cgtgaccata	tgttaaatgt	atggggctat	240
tctacaatga	acttcttttc	tccaatgagc	cggtatgctt	ccactgggtg	ggggccatta	300

gcagcttcat tagaatttaa gaaaatggtc aaggccttgc atagtgcagg aattgaggtt 360  
 attttggatg tggtttacia ccatacaaat gaagcggatg atgagcat 408

<210> 252

<211> 537

<212> DNA

<213> Eucalyptus grandis

<400> 252

ggcgaacacc agcacagtgt tatgctgact tcatgcgtgc atttagagac aacttccagc 60  
 accttttagg tgaaaccatt gtggaaattc aagtaggcat ggggccagca ggcgaacttc 120  
 gttatccatc ataccagag caaaatggga catggaaatt tccaggaatt ggagcttttc 180  
 aatgttacga caagtacatg ctgagtagct tgaaagctgc agccgaggct gctggcaaag 240  
 cagaatgggg ccacaccggt ccaactgatg ctggctacta taacaactgg ccggaggatg 300  
 cccatttctt caaaaaggaa ggtggaggat ggaacagtca atatggtgaa ttcttcttgt 360  
 cgtggtattc tcagatgcta ctggaccatg gtgagagaat actctcatct gccaaatcag 420  
 tctttgagaa tacaggaaca aagatttcag tcaagggtgc aggaatttca ctggcactat 480  
 ggaacgccgt tcgcatgctc ctgagctgac agcaggatac tacaacacac gttatcg 537

<210> 253

<211> 466

<212> DNA

<213> Eucalyptus grandis

<400> 253

gtagcttgaa agctgcagcc gaggtgctg gcaaagcaga atggggccac accggtccaa 60  
 ctgatgctgg tcaactataac aactggccag aggatgcccc attcttcaaa aagggaaggtg 120  
 gaggatggaa cagtcaatat ggtgaattct tcttgctgtg gtattctcag atgctactgg 180  
 accatggtga gagaatactc tcatctgccca aatcagtcct tgagaataca ggaacaaaga 240  
 tttcagtcga ggttgcagga attcaactggc actatggaac gcgttcgcat gctcctgagc 300  
 tgacagcagg atactacaac acacgttatc gggatgggta ccttcccatt gccagatgt 360  
 tggcacggca cgggtgctata ttcaacttca cttgcatcga aatgcgtgac cacgagcaac 420  
 cccaagatgc gctctgcgca cctgagaagc tgggtgaagca agtagc 466

<210> 254

<211> 364

<212> DNA

<213> Eucalyptus grandis

<400> 254

agatggcgaa gaagcatggg ttgaaagtgc aggtgtgtat gtcgtttcac cagtgcgggtg 60  
 gaaacgttgg tgactcttgc tccatccctc taccaaagtg ggctgtggaa gaagttgata 120  
 aagatccaga tcttgcatat acagaccagt ggggtaggag aaactacgag tacatatcgc 180  
 ttggctgtga caccctcccg gttctcaaag ggcgaacacc tgtacagtgt tatgctgact 240  
 tcatgcgtgc atttagagac aacttccagc accttttagg tgaaaccatt gtggaaattc 300  
 aagtaggcat ggggccagca ggcgaacttc gttatccatc ataccccag caaaatggga 360  
 catg 364

<210> 255

<211> 379

<212> DNA

<213> Eucalyptus grandis

<400> 255

ccagcatata cagaccagtg gggtaggaga aactacgagt acatatcgct tggctgggac 60  
 accctcccgg ttctcaaagg gcgaacacct gtacagtgtt atgctgactt catgcgtgca 120  
 ttttagagaca acttccagca ccttttaggt gaaaccattg tggaaattca agtaggcatg 180

gggccagcag	gcgaacttcg	ttatccatca	taccccgagc	aaaatgggac	atggaaattt	240
ccaggaattg	gagcttttca	atgttacgac	aagtacatgc	tgagtagctt	gaaagctgca	300
tccgaggctg	ctggcaaagc	agaatggggc	cacaccggtc	caactgatgc	tggtcactat	360
aacaactggc	cagaggatg					379

&lt;210&gt; 256

&lt;211&gt; 370

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 256

gaagctttcg	tgcagagttc	aatgactact	ttgaggatgg	tataatatca	atgattggga	60
ttggattggg	tccttgtggg	gagttacggg	acccatcaaa	ccctgtaaaa	aatgggttga	120
gatatcctgg	gataggtgaa	tttcagtgtc	acgatcagta	tctactgaag	aatctcagaa	180
aggcagcaga	ggcaaggggg	caggcttttt	gggctagagg	tccagataat	gcaggttctt	240
ataattcaca	gccacaagaa	actggtttct	tctgtgatgg	aggagattac	gatggctatt	300
ttggaagggt	cttccttaag	tggtactctc	aggtgttgat	tgatcatggg	gatagagtac	360
ttgccttggc						370

&lt;210&gt; 257

&lt;211&gt; 287

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 257

ggaaatctta	acaaggacat	ctactacaga	gatcggcatg	gatattctag	tgatgagtat	60
ctatctgctg	gagtggatca	aataacctata	ttatatggac	gtacagctgt	tgaatgctat	120
gaagatttca	tggtcagctt	catagacaaa	tttcaatcac	tcattgggaa	tccaattcaa	180
gaaattacta	ttggccttgg	tccgtcaggt	gaactaaggt	accctgcccc	tcctttttct	240
gatgggagat	ggaagttccc	tggtattgga	gaattccagt	gctatga		287

&lt;210&gt; 258

&lt;211&gt; 396

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 258

gcggtatcca	tcttatcctg	agagtaacgg	tacatggaaa	ttcccaggga	ttggagcatt	60
tcagtgtctat	gataagggtat	ggatactagc	catttttttt	tatgaatttc	cctccagctg	120
tatttttagtc	atatagttgc	ccttttttatt	tactagtgtg	gatttccttg	tattgcagta	180
tatgatctct	aaccttagat	ccacgtccga	agctgtctga	aagcaggagt	ggggtaattg	240
gggtccaagt	gatgcaggtc	attacaataa	ctggcctgag	gacagcccg	ttttccgcag	300
agatgggtgg	tggaacagtt	cttatgggtga	gttttttctt	gagtgggtatt	ctcgtatgct	360
tcttgatcat	ggagagagaa	tcctaggagc	agctga			396

&lt;210&gt; 259

&lt;211&gt; 420

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 259

gaggggtctca	tgcagcagaa	cttactgctg	ggtactataa	cacttcctat	agagatgggt	60
atgattcaat	cgctgcagtc	tttgcaagac	atggcgcagc	tttaaattat	ccttgtagtg	120
agatgttttg	tagtgaacag	ccagagatat	gctgctgcag	tccggagggt	ctcattaggc	180
agatgagaga	agttgcaagg	cgaggaaata	tacctttaac	aggtgaaaat	gcaattgaac	240
gctttgataa	ggaggctttc	tctcaaattg	tgagaaatgc	ttacaatcgt	cctcaggatg	300
tgagagcctt	tacgtatttc	cgaatgaggg	aggcactggt	caggactgat	aattggaaat	360

cattcgtgaa ctttggttaag cagatgtaca ataagtctca agatggaggc tgcaatggta 420

<210> 260

<211> 378

<212> DNA

<213> Pinus radiata

<400> 260

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gatacaatat ccgcttctaa cactttaaac cattgcaagg caattcaagc aggcttactt	120
gccttaaagg ctcttggtgt ggatggagtg gttatgcaag tattttgggg cattgtggag	180
agagatgctc caacaaaata tgactggtct gcataatttg ctttggtgaa aatgggtccaa	240
gcagcaggcc tgaaggttca ggcttcaata tgctttaatg gttgtaaatc tagtcaagaa	300
agcttgctca taccctctcc atcttggtt cttacagtgg gcaatagtga tccagacatc	360
ttcttcacag accgatcg	378

<210> 261

<211> 303

<212> DNA

<213> Pinus radiata

<400> 261

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tgggatgtga taatctccca gttttgaagg gaagaacgcc tgttcaatgc tatgtgatt	120
acatgaggag tttcaaggag aattttgggg atcttctggg agaaaccatt gtggaaattc	180
aagtgggaat gggtcctgca ggagagctga ggtatccatc ttatcctgag agtaatggta	240
cctggagatt cccaggaatt ggagcatttc agtgctatga taagtatatg gtctctaacc	300
tta	303

<210> 262

<211> 385

<212> DNA

<213> Pinus radiata

<400> 262

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ttttacatgc attgaaatgc aggataatga acaaccgtcc gaagccagtt gtgggcccga	120
ggctcttggt cgtcaggat taaatgctgg atggaaagaa ggtattgagg ttctctgtga	180
aaatgcttta cctagatttg atgaagaagc atatgatcag attgtaaggc aatccagacc	240
cgagggaata aacgaaacag gaccgcccga gaagcgaata tctgctttta cctatctaag	300
gctatctcaa gaactcatgc aagaacatag ctggaaagaa ttcaacaaat tcctgagaag	360
aatgcacgtg agtttggatt atcat	385

<210> 263

<211> 330

<212> DNA

<213> Pinus radiata

<400> 263

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tgccagtggg gtgacggctc cgagcccacg ggtaactggg ccccatgaa cctgggtgtt	120
ggctactctg ccggtccac ctggttgctc atgttcaaga acgagcctac cactctgtcc	180
aacctggact tcaagatcaa gattatcggt gacgacctga gcgacaactg cagctacgac	240
ggagccggca acttctacaa caaggccggc ctgatcacct ctggcaacgg ctgcaactgc	300
agctcctcct ctggcaacgc ctacttcgtc	330

<210> 264



<211> 359  
 <212> DNA  
 <213> Eucalyptus grandis

<400> 264  
 aatttggaca tcctgaatgg ttggactttc caagggaagg gaacgggttc agttatgata 60  
 aatgccggcg aagatttgat ctaggagatg ctgagtatct aagataccgg ggaatgcaag 120  
 aattttgatca ggctatgcag catgttgaag aagcttatgg cttcatgact tctgagcacc 180  
 aatacatatc cagaaaagat gaaggagaca gggcatcgt ctttgaaagg ggaaatcttg 240  
 tgtttgtctt caatttccat tggaaataata gctacacgga ctaccatgta ggctgcttga 300  
 agcctggaaa gntaagattg tcttaaattc agatgacgcc ttgtttggag ggtatagta 359

<210> 265  
 <211> 451  
 <212> DNA  
 <213> Pinus radiata

<400> 265  
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 aagccaatgc tgtgcctatc caggccatcg aagaacaaac tactaagcca atgcaaaaca 120  
 gagttgagct tgaagatcgg ccaaaagtgg tccctccacc tgggagtggc caaaggattt 180  
 atgaaataga tccattgttg aataactatc gtgaacatct tgattatcga tttgcgagct 240  
 ataagaaaac aagagagttg attgataaat atgaagggtg cttggaagca ttttctaggg 300  
 gttatgaaaa gatgggattt aatagaagtg cggctggaat cacatacaga gagtgggcac 360  
 ctggtgctaa gggggcatca cttataggag atttcaacaa ctggaatccc aatgctgatg 420  
 ttatgctaag aacgagtttg gagtatggga g 451

<210> 266  
 <211> 375  
 <212> DNA  
 <213> Pinus radiata

<400> 266  
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 gggagatatt ttaccacaaac aatgcagatg gttctccacc aattcccat ggatctcgtg 120  
 ttaagataca tatggataca atttcgggac caaaggacgc aatccctgct tggattaagt 180  
 ttgctgttca agctccgggt gagattccat acaatggaat atattatgat cccctcctg 240  
 aggataaata tgaatttaag tatcctcgac caaagcagcc caaatcattg cgaatatatg 300  
 aagcgcatgt tggcatgagc agcacggaac ccaaaattaa tacatatgtt gagttcaggg 360  
 atgatgtact accac 375

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 <211> 408  
 <212> DNA  
 <213> Pinus radiata

<400> 267  
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 caaagcaaath agtttagcatt gcaaatgaag aagataaggt gattgttttt gaacgaggcg 360  
 atatagtgtt tgttttcaac ttccatccca agaatacata tcctgggt 408

<210> 268  
 <211> 476

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 268

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agtacttcca tggaggacca aagggatacc acgacttggt ggacagtcgt ctcttcaact	360
actctcatta tgaggttctc cgcttcctga tgtctaactc gcgcttctgg atggaagagt	420
atcagtttga cggtttccgc tttgatgggtg ttacgagtat gttgtatctg catcac	476

&lt;210&gt; 269

&lt;211&gt; 313

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 269

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tgggaccgac actctcatgc tatgacaaa gatcagtatg gtgtctgggg aatcacgatc	180
cctagcatca acggacagcc tgctatcccc cacgattcga aaataaagggt ttcgttcggt	240
attcctggcg gcgagcgtat cgagcgtctg cctgcttgga tcaagcgcgt caccaggagc	300
ctctctgtct cgc	313

&lt;210&gt; 270

&lt;211&gt; 258

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 270

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gattcatggg ctatttcccc aggcataaac aattggagaa gatgtgagtg gtatgccaac	180
cttttctcgt tctgtacaag atggcggagt gggatttgat tatcgactcc atatggcagt	240
agctgacaaa tggattga	258

&lt;210&gt; 271

&lt;211&gt; 349

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 271

cctatccagg ctttcgaaga acaaactact aagccaatgc aaaacagagt ngagcttgaa	60
gatcggccaa aagttgtccc tccacctggg agtggccaaa ggatttatga aatagatcca	120
ttgttgaata actatcgtga acatcttgat tatcgatttg cgcagtataa gaaaacaaga	180
gagttgattg ataaatatga aggtggcttg gaagcatttt ctaggggtta tgaaaagatg	240
ggattttaata gaagtgcggc tggaatcaca tacagagagt gggcacctgg tgctaagggg	300
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&lt;210&gt; 272

&lt;211&gt; 369

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 272

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cgcaacatgt	ttgtatgact	gaagaaagca	gtattttctaa	aacagatatc	ggaggacatg	300
gcttctccca	ttatcttgtt	tttccatata	aatctcactt	gggcatacca	tcttttagttc	360
tgtagtcag						369

&lt;210&gt; 273

&lt;211&gt; 327

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 273

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cgatcgccgt	gacccgcctc	ggcgcccggt	ccgcgttcgt	cggcaaagct	cggggacgac	300
gagttcgggc	aacaatgctg	gccggat				327

&lt;210&gt; 274

&lt;211&gt; 275

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 274

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ccaagtgcgg	ccatgctttt	acatgaatcg	gaaactagat	gttgaactta	tcagcaaggc	180
aaagatcttc	cattatgggt	ctatcagttt	gattgatgaa	ccctgcaaat	cggctcatct	240
tcagcaaatg	aaaattgcca	aaaactcagg	gagcg			275

&lt;210&gt; 275

&lt;211&gt; 362

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 275

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cccggcgggc	cccccgcaac	gtcgcgatcg	ccgtgaccgc	cctcggcggc	cgggtccgagt	180
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taacgctccg	tgccgacggg	gagcgcgagt	tcatgttcta	cgggaaccgc	agcgcggaca	360
tg						362

&lt;210&gt; 276

&lt;211&gt; 543

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 276

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taaccagact	ctcttcaatt	gaaaatggaa	gcaagcaacg	actagtccta	agggaaaatc	180
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274  
12/0  
10

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cat						543

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<212> DNA  
<213> Eucalyptus grandis

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<210> 278  
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<213> Eucalyptus grandis

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<212> DNA  
<213> Eucalyptus grandis

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acgcgtgcgg	agccatcacc	accaccaaga	agggagcgat	ccccgccctc	ccgaccgagg	180
ctgatgtcct	cagcttgatt	g				201

<210> 280  
<211> 319  
<212> DNA  
<213> Eucalyptus grandis

<400> 280						
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acgctggggc	tctgctctcc	tatgatccaa	acctcagact	accattgtgg	ccatcacctg	240
aggaggctcg	tgagcagatc	aagagcattt	ggggacaaag	gcagatatca	tttaaaagtg	300
gagtgattgt	tgaactgga					319

<210> 281  
<211> 446  
<212> DNA  
<213> Eucalyptus grandis

&lt;400&gt; 281

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ccggaacccg	agcgccgaca	tgctgctcaa	gcccaggagg	ctcaacctcg	agctgattan	420
gatctgcgaa	agtctttcat	tatgga				446

&lt;210&gt; 282

&lt;211&gt; 369

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 282

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acctgaggag	gctcgtgagc	agatcaagag	catttggggac	aaggcagata	tcattaaagt	360
gagtgatgt						369

&lt;210&gt; 283

&lt;211&gt; 583

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 283

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ccatcttgaa	gcaatgcaag	ttgccaagga	cgctggggct	ctgctctcct	atgatccaaa	540
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&lt;210&gt; 284

&lt;211&gt; 305

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 284

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ggggacgacg	agttcgggca	catgctggcc	gggatcctga	aaggagaacg	gggtcaactg	300
cgacg						305

&lt;210&gt; 285

&lt;211&gt; 403

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 285

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&lt;210&gt; 286

&lt;211&gt; 471

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 286

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actaccattg	tgggcatcac	ctgaggaggt	tcgtgagcag	atcaagagca	t	471

&lt;210&gt; 287

&lt;211&gt; 410

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 287

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&lt;210&gt; 288

&lt;211&gt; 451

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 288

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 <213> Eucalyptus grandis

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<210> 290  
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 <212> DNA  
 <213> Eucalyptus grandis

<400> 290  
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<210> 291  
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 <212> DNA  
 <213> Eucalyptus grandis

<400> 291  
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 gaggcccccg ggttcctcaa gggccccggc ggccgccccg ccaacgtcgc gatcgccgtg 180  
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 <212> DNA  
 <213> Eucalyptus grandis

<400> 292  
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<210> 293  
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 <212> DNA  
 <213> *Eucalyptus grandis*

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<210> 294  
 <211> 329  
 <212> DNA  
 <213> *Pinus radiata*

<400> 294  
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<210> 295  
 <211> 496  
 <212> DNA  
 <213> *Pinus radiata*

<400> 295  
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 gttggggatg atgagtttgg gcgcatgctt gctgacattc tgagggaaaa caatgtgatg 300  
 gaccgaggaa ttagatttga ttcccatgcc agaaccgcgc tggcattcgt tactttaaag 360  
 atgaatggcg agagggaatt tatgttctat cgtaatccca gcgctgacat gcttctcaag 420  
 gaatctgagc ttgatgcaga gctgatccga gaggcacga tatttcacta tggatcaatc 480  
 agtctgattg cagagc 496

<210> 296  
 <211> 473  
 <212> DNA  
 <213> *Pinus radiata*

<400> 296  
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 ggtgcacctg ctaatgttgc tgtttgcata gcaagactag gaggttcac agcatttatt 180  
 ggaaagggtg gtgatgacga gtttgatgat atgcttgctg atatcttggg gaaaaacaat 240  
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 ttaaggagtg atggtgaacg tgaatttatg ttttacagaa atccaagtgc agatatgtta 360  
 ctcgacgaat cagagcttga tgtggatctc atcagagagg caaaaatttt ccactatggt 420  
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<210> 297



<211> 369  
 <212> DNA  
 <213> Pinus radiata

<400> 297  
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 gcacctgcta atgttgctgt ttgcatagca agactaggag gttcatcagc atttattgga 180  
 aagggttggtg atgacgagtt tggatatatg cttgctgata tcttgaggaa aaacaatgta 240  
 aataataagg gcatgcgttt tgatgctgga gctcgaactg ctttggcatt tgtgacatta 300  
 aggagtgatg gtgaacgtga atttatgttt tacagaaatc caagtgcaga tatgttactc 360  
 gacgaatca 369

<210> 298  
 <211> 459  
 <212> DNA  
 <213> Pinus radiata

<400> 298  
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<210> 299  
 <211> 417  
 <212> DNA  
 <213> Pinus radiata

<400> 299  
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<210> 300  
 <211> 359  
 <212> DNA  
 <213> Pinus radiata

<400> 300  
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 tcgttcacca tcggctttca atgtgacaaa agccaaagca gttcgagcac caagatcaaa 180  
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<210> 301  
 <211> 374

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 301

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cactccattg	accg					374

&lt;210&gt; 302

&lt;211&gt; 339

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 302

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&lt;210&gt; 303

&lt;211&gt; 402

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 303

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&lt;210&gt; 304

&lt;211&gt; 468

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 304

gcgaatgttg	ccgttggtat	agcaaggctc	ggaggttcat	ccgcatttat	agggaagggtt	60
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ctgatcacgg	accctgttaa	gtctgcgcac	ttggctgcc	tgaaaatcgc	tagagacacg	360
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&lt;210&gt; 305

&lt;211&gt; 502

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 305

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&lt;210&gt; 306

&lt;211&gt; 379

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 306

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atccttggtga	tgatgctgt					379

&lt;210&gt; 307

&lt;211&gt; 233

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 307

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&lt;210&gt; 308

&lt;211&gt; 377

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 308

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&lt;210&gt; 309

&lt;211&gt; 517

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

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<210> 310  
 <211> 360  
 <212> DNA  
 <213> Pinus radiata

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<210> 311  
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 <212> DNA  
 <213> Pinus radiata

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<210> 312  
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 <213> Pinus radiata

<400> 312  
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<210> 313  
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 <212> DNA  
 <213> Pinus radiata

<400> 313

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&lt;210&gt; 314

&lt;211&gt; 487

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 314

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&lt;210&gt; 314

&lt;211&gt; 421

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 315

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t						421

&lt;210&gt; 316

&lt;211&gt; 420

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 316

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&lt;210&gt; 317

&lt;211&gt; 499

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 317

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&lt;210&gt; 318

&lt;211&gt; 364

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 318

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ttac						364

&lt;210&gt; 319

&lt;211&gt; 298

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 319

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&lt;210&gt; 320

&lt;211&gt; 261

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 320

gttttcgaga	gagaaatggc	gaaggcgatt	cgtttcttcg	agctcaaacac	cgggggccaag	60
atcccctcgc	tcgggctggg	cacttggcag	accggtgacg	gcgtcgacgc	cgtcaccacc	120
gccatcaagg	ttgggtacag	gcatattgat	tgtgctcaag	cttatcaaaa	tgagaaggag	180
attgggtactg	ctctccagaa	attattcagc	gagggtgtgg	tgaagcgcga	ggatttgtgg	240
atcacatcca	agctatggtg	t				261

&lt;210&gt; 321

&lt;211&gt; 450

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

## &lt;400&gt; 321

ctaggggtgag atcgaagctc gcgttttcga gagagaaatg gcgaaggcga ttcgtttctt	60
cgagctcaac accggggcca agatcccctc cgtcgggctg ggcacttggc agaccggtga	120
cggcgtcgac gccgtcacca ccgccatcaa ggttgggtac aggcattatg attgtgctca	180
agcttatcaa aatgagaagg agattggtac tgctctccag aaattattca gcgaggggtg	240
ggtgaagcgc gaggatttgt ggatcacatc caagctatgg tgtgctgac acgcaccaga	300
agatgttccc aaggcattag aaagaaccct ggagaacttg cagctcgagt atctggatct	360
ttacctgac cactggccgg tgagcatgag gaaggctcaa tttggcttca agcctgaaaa	420
ccttaccag ccggacatac ccagtacgtg	450

## &lt;210&gt; 322

## &lt;211&gt; 347

## &lt;212&gt; DNA

## &lt;213&gt; Eucalyptus grandis

## &lt;400&gt; 322

cgagagagaa atggcgaagg cgattcgttt cttcgagctc aacaccgggg ccaagatccc	60
ctccgtcggg ctgggcactt ggcagaccgg tgacggcgctc gacgcggtca ccaccgccat	120
caaggttggg tacaggcata ttgattgtgc tcaagcttat caaaatgaga aggagattgg	180
tactgctctc cagaaattat tcagcgaggg tgtggtgaag cgcgaggatt tgtggatcac	240
atccaagcta tgggtgtgctg atcacgcacc agaagatggt cccaaggcat tagaaagaac	300
cctggagaac ttgcagctcg agtatctgga tctttacctg atccact	347

## &lt;210&gt; 323

## &lt;211&gt; 414

## &lt;212&gt; DNA

## &lt;213&gt; Eucalyptus grandis

## &lt;400&gt; 323

cacgagtcgg ctggcgaagt agtcgaagta ggtgaaggcg taaccaaatg gaaagtcggc	60
gaccgagtcg ctatcgaggc tggagtacct tgttcccaac ctgcttgcca tgcgtgtcgt	120
actggccgat acaacgcacg ccagatgtc gttttcttct caaccccgcc gttccatggc	180
acattgacgc gctggcacct tcatccggca cagtgggtgc accgtcttcc ggataatggt	240
tctttcgaag agggcgccct gtgcgaacca ctgctgtgc cattggccgg catcgagcgt	300
tccggtctca gactcggaga tcccgctcct gtctgtggtg ctggaccaat aggcctaata	360
tctctacttt cggcccgtgc tgcgggtgca gagcctattg ttataacgga cctt	414

## &lt;210&gt; 324

## &lt;211&gt; 464

## &lt;212&gt; DNA

## &lt;213&gt; Eucalyptus grandis

## &lt;400&gt; 324

cacgagtcgg ctggcgaagt agtcgaagta ggtgaaggcg taaccaaatg gaaagtcggc	60
gaccgagtcg ctatcgaggc tggagtacct tgttcccaac ctgcttgcca tgcgtgtcgt	120
actggccgat acaacgcacg ccagatgtc gttttcttct caaccccgcc gttccatggc	180
acattgacgc gctggcacct tcatccggca cagtgggtgc accgtcttcc ggataatggt	240
tctttcgaag agggcgccct gtgcgaacca ctgctgtgc cattggccgg catcgagcgt	300
tccggtctca gactcggaga tcccgctcct gtctgtggtg ctggaccaat aggcctaata	360
tctctacttt cggcccgtgc tgcgggtgca gagcctattg ttataacgga ccttttccaa	420
agccgtctgg actttgcgaa gaagctggtg cctggcgctc gcac	464

## &lt;210&gt; 325

## &lt;211&gt; 368

## &lt;212&gt; DNA

## &lt;213&gt; Eucalyptus grandis

&lt;400&gt; 325

cacgagtcgg	ctggcgaagt	agtcgaagta	ggtgaaggcg	taacccaatg	gaaagtcggc	60
gaccgagtcg	ctatcgaggc	tggagtacct	tgttcccaac	ctgcttgcca	tgcgtgtcgt	120
actggccgat	acaacgcatg	cccagatgtc	gttttcttct	caaccccgcc	gttccatggc	180
acattgacgc	gctggcacct	tcatccggca	cagtgggtgc	accgtcttcc	ggataatgtt	240
tctttcgaag	agggcgccct	gtgcgaacca	ctcgctgtcg	cattggccgg	catcgagcgt	300
tccggtctca	gactcggaga	tcccgtcctt	gtctgtggtg	ctggaccaat	aggcctaata	360
tctctact						368

&lt;210&gt; 326

&lt;211&gt; 350

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 326

ctaggggtgag	atcgaagctc	gcgttttcga	gagagaaatg	gcgaaggcga	ttcgtttctt	60
cgagctcaac	accggggcca	agatccccct	cgtcgggctg	ggcacttggc	agaccgggtga	120
cggcgctcgac	gccgtcacca	ccgccatcaa	ggttgggtac	aggcatattg	atttgtgtca	180
agcttatcaa	aatgagaagg	agattggtac	tgtctctccag	aaattattca	gcgaggggtgt	240
ggtgaagcgc	gaggatttgt	ggatcacatc	caagctatgg	tgtgctgata	acgcaccaga	300
agatgttccc	aaggcattag	aaagaaccct	ggagaacttg	cagctcgagt		350

&lt;210&gt; 327

&lt;211&gt; 372

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 327

cttcgagctc	aacaccgggg	ccaagatccc	ctccgtcggg	ctgggcactt	ggcagaccgg	60
tgacggcgctc	gacgccgtca	ccaccgccat	caagggtggg	tacaggcata	ttgattgtgc	120
tcaagcttat	caaaatgaga	aggagattgg	tactgctctc	cagaaattat	tcagcgaggg	180
tgtggtgaag	cgcgaggatt	tgtggatcac	atccaagcta	tgggtgtgctg	atcacgcacc	240
agaagatgtt	cccaaggcat	tagaaagaac	cctggagaac	ttgcagctcg	agtatctgga	300
tctttacctg	atccactggc	cggtgagcat	gaggaagggc	tcaattgggt	tcaagcctga	360
aaaccttacc	ca					372

&lt;210&gt; 328

&lt;211&gt; 333

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 328

aaatggcgaa	ggcgattcgt	ttcttcgagc	tcaacaccgg	ggccaagatc	ccctccgtcg	60
ggctgggcac	ttggcagacc	ggtgacggcg	tcgacgccgt	caccaccgcc	atcaagggtg	120
ggtacaggca	tattgattgt	gctcaagctt	atcaaaatga	gaaggagatt	ggtactgtctc	180
tccagaaatt	attcagcgag	ggtgtggtga	agcgcgagga	tttgtggatc	acatccaagc	240
tatggtgtgc	tgatcacgca	ccagaagatg	ttcccaaggc	attagaaaga	accctggaga	300
acttgcagct	cgagtatctg	gatctttacc	tga			333

&lt;210&gt; 329

&lt;211&gt; 377

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 329

cgcaactcctt	ttgcctgccc	cccacgagcc	cggagcggga	gtagactgag	atcgaagctc	60
gcgttttcga	gagagaaatg	gcgaaggcga	ttcgtttctt	cgagctcaac	acggggggcca	120



agatccccctc	catcgggctg	ggcacttgge	aggccgatcc	cggcgtcgtg	gccgaggccg	180
tcaccaccgc	cacaaaggct	gggtacagge	atattgattg	tgctcaagct	tattacaatg	240
agaaggagat	tggtagctg	ctccagaaat	tattcagcga	gggtgtgggtg	aagcgcgagg	300
atttgtggat	cacttccaag	ctatggtgta	ctgatcacgc	accggaagat	gttcccaagg	360
caatagacag	aaccttg					377

&lt;210&gt; 330

&lt;211&gt; 484

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 330

aactaatggt	tcgcttgcca	ataagggtggt	gcatcctgca	catctgtggt	acaagctacc	60
ggaaaatgtg	agcttggagg	aaggagcaat	gtgtgaaccc	ctcagtgttg	gtgtacacgc	120
ttgtcgccga	gcaaatatca	atcctgagac	caacatactc	ataataggat	cagggccgat	180
tggccttggt	accttattag	cagcccggtg	ttttggagct	ccgagaatcg	tcatcactga	240
tgtagacgag	tgcagattat	cgattgcaaa	aatgcttggt	gcctctgagg	tggttcaagt	300
ctcaacagat	gttcagctag	tggatgaaga	agtggcgcg	atccaaaatg	caatgggctg	360
cgacattgat	gtgagcttcg	attgtgttgg	ctatgacaag	acaatgacca	cagctttgaa	420
tgcgactcgt	gctgggtggca	aagtgtgcct	catcggacta	gccttgagca	agatgacagt	480
tcct						484

&lt;210&gt; 331

&lt;211&gt; 477

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 331

ccaaagggaa	aaaaaatggg	gaagggagca	atgtctcagg	gtaacgaaaa	tggggaaggt	60
gacaatatgg	ctgcatggct	cactggaata	aacactcttc	gcattccagcc	cttcaaactt	120
ccgcctcttg	gcccccatga	tgcgaagggtg	cgcatgaagg	ctgtgggtat	ctgtggcagt	180
gacgtccact	atttgaggac	attacgggtg	gcggacttta	ttgtaaaaga	gccaatgggtg	240
attggctcatg	agtctgctgg	aataattgag	gaggttggca	gtgaagtga	acatctgggt	300
cctgggtgacc	gcgtactttg	gagcctggaa	tatcgtgttg	gcgttgtgac	caatgtaagc	360
gaggctccta	caatttgtgt	cccagatga	agttttttgc	aacacctccc	gtgcatgggt	420
ccttggtccaa	tcagattggt	catcctgcag	atttatgttt	caagttgcca	gataatg	477

&lt;210&gt; 332

&lt;211&gt; 433

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 332

agggtaacga	aaatggggaa	ggtgacaata	tggctgcatg	gctcactgga	ataaacactc	60
ttcgcaccca	gcccttcaaa	cttcgcctc	ttggccccc	tgatgcgaag	gtgcgcgatga	120
nggctgtggg	tatctgtggc	agtgaagctc	actatttgan	gacattacgg	tgtgcggact	180
ttattgtaaa	anagccaatg	gtgattgggtc	atgagctctg	tggaataatt	gaggagggtg	240
gcagtgaagt	gaaacatctg	gttcctgggtg	accgcgtact	ttggagcctg	gaatatcgtg	300
ttggcgttgt	gaccaatgta	agcnaggctc	ctacaatttg	tgccccgaga	tgaagttttt	360
tgcaacacct	cccgtgcatg	gttccttgge	caatcagatt	gttcacctctg	cagatttatg	420
tttcaagttg	cca					433

&lt;210&gt; 333

&lt;211&gt; 466

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 333

gaggaatagg	aacaggaccc	tcttttggtg	ggcacacttg	catctgtctt	tgttccagca	60
atgtcaggag	cagctgctgc	gattacactg	aataatggcc	acaaaatgcc	catcattggg	120
cttgagtggt	ggagaatgga	gggtcaggaa	ataagagacc	ttatcttcaa	tgcactacac	180
ataggggtacc	gtcatttcga	ttgtgcagct	gattacagga	atgaaaagga	agttgggtcaa	240
gcacttgccg	aggcctttca	gcaaggcttg	gtgaaacgag	aggatatttt	tattactacc	300
aagctatgga	attcagacca	tggacatggt	cttgaggcat	gcaaggacag	tttaaagaat	360
ctgcagttgg	aatatttgga	cctgtacttg	gttcattttc	caatagccac	acgacataca	420
ggggttggaa	caactgatag	tgccttagac	gaagatggtg	ttctcg		466

&lt;210&gt; 334

&lt;211&gt; 483

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 334

gggtaacgaa	aatggggaag	gtgacaatat	ggctgcatgg	ctcactggaa	taaacactct	60
tcgcatccag	cccttcaaac	ttccgcctct	tggcccccac	gatgcgaagg	tgcgcatgaa	120
ggctgtgggt	atctgtggca	gtgacgtcca	ctatttgagg	acattacggt	gtgcggactt	180
tattgtaaaa	gagccaatgg	tgattgggtca	tgagtctgct	ggaataattg	aggaggttgg	240
cagtgaagtg	aaacatctgg	ttcctgggtga	ccgcgtagct	ttggagcctg	gaatatcgtg	300
ttggcggtgt	gaccaatgta	agcgaggctc	ctacaatttg	tgtcccgaga	tgaagttttt	360
tgcaacacct	cccgtgcatg	gttccttgge	caatcagatt	gttcatcctg	cagatttatg	420
tttcaagttg	ccagataatg	taagtctcga	ggaagggtgcc	atgtgtgaac	cactcagtgt	480
tgg						483

&lt;210&gt; 335

&lt;211&gt; 329

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 335

gggtaacgaa	aatggggaag	gtgacaatat	ggctgcatgg	ctcactggaa	taaacactct	60
tcgcatccag	cccttcaaac	ttccgcctct	tggcccccac	gatgcgaagg	tgcgcatgaa	120
ggctgtgggt	atctgtggca	gngacgtcca	ctatttgagg	acattacggt	gtgcggactt	180
tattgtaaaa	gagccaatgg	tgattgggtca	tgagtctgct	ggaataattg	aggaggttgg	240
cagtgaagtg	aaacatctgg	ttcctgggtga	ccgcgtagct	ttggagcctg	gaatatcngn	300
ttggcggtgn	gaccaatgta	agcgaggctt				329

&lt;210&gt; 336

&lt;211&gt; 419

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 336

ctcagggtaa	cgaaaatggg	gaagggtgaca	atatggctgc	atgggtcact	ggaataaaca	60
ctcttcgcat	ccagcccttc	aaacttccgc	ctcttgcccc	ccatgatgcg	aagggtgcga	120
tgaaggctgt	gggtatctgt	ggcagtgacg	tccactattt	gaggacatta	cgggtgtgcg	180
actttattgt	aaaagagcca	atgggtgattg	gtcatgagtc	tgctggaata	attgaggagg	240
ttggcagtg	agtgaacat	ctgggttcctg	gtgaccgcgt	agctttggag	cctggaatat	300
cgtgttggcg	ttgtgaccaa	tgtaagcgag	gctcctacaa	tttgtgtccc	gagatgaagt	360
tttttgcaac	acctcccgtg	catgggttctt	tggccaatca	gattgttcat	cctgcagat	419

&lt;210&gt; 337

&lt;211&gt; 392

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 337

ctcagggtaa	cgaaaaatggg	gaaggtgaca	atatggctgc	atggctcact	ggaataaaca	60
ctcttcgcat	ccagcccttc	aaacttccgc	ctcttgcccc	ccatgatgcg	aaggtgcgca	120
tgaaggctgt	gggtatctgt	ggcagtgacg	tccactatct	gaggacatta	cgggtgcgcg	180
actttattgt	aaaagagcca	atgggtgattg	gtcatgagtc	tgctggaata	attgaggagg	240
ttggcagtg	agtgaacat	ctggttcctg	gtgaccgcgt	agctttggag	cctggaatat	300
cgtgttggcg	ttgtgaccaa	tgtaagcgag	gctcctacaa	tttgtgtccc	gagatgaagt	360
tttttgcaac	acctcccgtg	catggttcct	tg			392

&lt;210&gt; 338

&lt;211&gt; 362

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 338

ctaaccaaaag	ggaaaaaaaa	tggggaaggg	agcaatgtct	cagggtaacg	aaaatgggga	60
aggtgacaat	atggctgcat	ggctcactgg	aataaacact	cttcgcatcc	agcccttcaa	120
acttcgcct	cttgccccc	atgatgcgaa	ggtgcgcatg	aaggctgtgg	gtatctgtgg	180
cagtgcagtc	cactatctga	ggacattacg	gtgtgcggac	tttattgtaa	aagagccaat	240
ggtgattggt	catgagtcgt	ctggaataat	tgaggagggt	ggcagtgaag	tgaaacatct	300
ggttcctggt	gaccgcgtag	ctttggagcc	tggaatatcg	tggtggccgt	tgtgaccaat	360
gt						362

&lt;210&gt; 339

&lt;211&gt; 417

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 339

aaaaaatgg	ggaagggagc	aatgtctcag	ggtaacgaaa	atgggggaagg	tgacaatatg	60
gctgcatggc	tcactggaat	aaacactctt	cgcatccagc	ccttcaaact	tccgcctctt	120
ggcccccag	atgcgaagg	gcgcaggaag	gctgtgggta	tctgtggcag	tgacgtccac	180
tatttgagga	cattacgggt	tgccgacttt	attgtaaaag	agccaatgg	gattgggtcat	240
gagtctgctg	gaataattga	ggaggttggc	agtgaagtga	aacatctggt	tcctgggtgac	300
cgcgtagctt	tgagccctgg	aatatcgtgt	tgccgtttgt	accaatgtaa	gcgaggctcc	360
tacaatttgt	gtcccagat	gaagtttttt	gcaacacctc	ccgtgcatgg	ttccttg	417

&lt;210&gt; 340

&lt;211&gt; 343

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 340

ccaaagggaa	aaaaatgggg	aaggagcaa	tgtctcaggg	taacgaaaat	ggggaagggtg	60
acaatatggc	tgcatggctc	actggaataa	acactcttcg	catccagccc	ttcaaacttc	120
cgctctcttg	cccccatgat	gcgaagggtg	gcatgaaggc	tgtgggtatc	tgtggcagtg	180
acgtccacta	tttgaggaca	ttacgggtgtg	cggactttat	tgtaaaagag	ccaatgggtga	240
ttggtcatga	gtctgctgga	ataattgagg	agggttggcag	tgaagtga	catctgggttc	300
ctgggtgaccg	cgtagctttg	gagcctggaa	tatcgtgttg	gcg		343

&lt;210&gt; 341

&lt;211&gt; 590

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 341

attggtcatg	agtctgctgg	aataattgag	gaggttgcca	gtgaagtga	acatctgggt	60
cctggtgacc	gcgtagcttt	ggagcctgga	atatacgtgtt	ggcgttgtga	ccaatgtaag	120
cgaggctcct	acaattttgtg	tcccagatg	aagttttttg	caacacctcc	cgtgcatggg	180
tccttgccca	atcagattgt	tcatacctgca	gatttatgtt	tcaagttgcc	agataatgta	240
agctcgcagg	aagggtcccat	gtgtgaacca	ctcagtggtg	gggttcacgc	ttgtcgccgt	300
gcttctgtag	gccctgagac	aaatgtcttg	gtaatggggg	caggtccctat	cggccttgct	360
accgtgctgt	ctgcacgtgc	atttgagct	tcacgaatta	ttattgctga	tgtagatgaa	420
gagcgtctgt	caatggctaa	aaaggttggc	tccgatgaat	gcgtcttagt	ctccagagac	480
tctcaggata	ttgatgaaga	agtgacccgc	atacaaaatg	ccatgggtgg	aaacatagat	540
gtaacttttg	attgtgctgg	ttttgctaaa	accatgtcga	cggctctaaa		590

&lt;210&gt; 342

&lt;211&gt; 372

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 342

atctaacc	aaaggga	aaaaaa	atgggga	agg	gagcaat	gtc	tcagggt	aaac	gaaaa	gggg	60	
aagggtg	acaa	tatggct	gca	tggctc	actg	gaataa	acac	tcttcg	catc	cagccct	tca	120
aacttcc	gcc	tcttg	cccc	catgat	gcga	aggtgc	gcgat	gaaggc	gtg	ggatat	ctgtg	180
gcagtga	cgt	ccactat	ttg	aggacat	tac	ggtgtg	cgga	ctttatt	gta	aaagag	ccaa	240
tggtgat	tgg	tcatga	gtct	gctgga	ataa	ttgagg	agg	tggcag	tga	gtgaa	acatc	300
tggttcc	tgg	tgaccg	cgt	gctttg	gagc	ctggaat	atc	gtgttg	ggcg	tgtgac	caat	360
gtaagcg	agg	ct										372

&lt;210&gt; 343

&lt;211&gt; 378

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 343

gtggcag	tga	cgtcc	actat	ttgagg	acat	tacgg	tgtgc	ggact	tttatt	gtaaa	agagc	60
caatgg	tgat	tggtc	atgag	tctgt	cgaa	taatt	gagga	ggttg	gcag	gaagt	gaaac	120
atctgg	ttcc	tggtg	accgc	gtagc	tttg	agcct	ggaat	atcgt	gttgg	cgttg	tgacc	180
aatgta	agcg	aggct	cctac	aattt	gtgtc	ccgag	atgaa	gtttt	ttgca	acacc	tccc	240
tgcatg	gttc	cttgg	ccaat	cagatt	gttc	atcct	gcaga	tttat	gtttc	aagtt	gccag	300
ataatg	taag	tctcg	aggaa	ggtgc	catgt	gtga	accact	cagt	gttggg	gttcat	gttc	360
gtcgccc	gtg	cttct	gtga									378

&lt;210&gt; 344

&lt;211&gt; 510

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 344

agcaatg	tct	cagggt	aaacg	aaaat	ggggg	aggtg	acaat	atggg	ctgcat	ggctc	actgg	60
aataaac	act	cttcg	catcc	agcc	cttcaa	actt	ccgcct	cttgg	cccc	atgat	gcgaa	120
ggtgcg	catg	aagg	ctgtg	gtat	ctgtg	cagt	gacgtc	cact	atttga	ggac	attacg	180
gtgtgc	ggac	tttatt	gtaa	aagag	ccaat	ggtg	attgg	catg	agtctg	ctgga	ataat	240
tgagga	aggt	ggcag	tgaag	tga	acatct	ggtt	cctgg	gacc	gcgtag	ctttg	ggagcc	300
tggaat	atcg	tggt	ggcgt	gtg	accaatg	taag	cgaggc	tct	tacaat	tgtg	tcccg	360
gatga	agttt	tttg	caacac	ctccc	gtgca	tggt	tcctg	gcc	aatcaga	ttgt	tcatcc	420
tgca	gattta	tg	tttca	agt	tgcc	agataa	tg	taagt	ctc	gagga	aggtg	480
accact	cagt	g	ttggg	gttc	atg	ctt	gtc					510

&lt;210&gt; 345

&lt;211&gt; 504

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 345

gtctcagggt	aacgaaaatg	gggaagggtga	caatatggct	gcatggctca	ctggaataaa	60
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catgaaggct	gtgggtatct	gtggcagtga	cgtccactat	ttgaggacat	tacgggtgtgc	180
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atcgtgttgg	cgttgtgacc	aatgtaagcg	aggctcctac	aatttgtgtc	ccgagatgaa	360
gttttttgca	acacctcccg	tgcattggttc	cttggccaat	cagattgttc	atcctgcaga	420
tttatgtttc	aagttgccag	ataatgtaag	tctcgaggaa	ggtgccatgt	gtgaaccact	480
cagtgttggg	gttcatgttc	gtcg				504

&lt;210&gt; 346

&lt;211&gt; 426

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 346

gcaatgtctc	agggtaacga	aaatggggaa	ggtgacaata	tggtgcatg	gctcactgga	60
ataaacactc	ttcgcatcca	gcccttcaaa	cttcgcctc	ttggcccca	tgatgcgaag	120
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atgaagtttt	ttgcaacacc	tcccgtgcat	ggttccttgg	ccaatcagat	tgttcatcct	420
gcagat						426

&lt;210&gt; 347

&lt;211&gt; 534

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 347

atattgtgtcc	cgagatgaag	ttttttgcaa	cacctcccgt	gcatgggttcc	ttggccaatc	60
agattgttca	tcctgcagat	ttatgtttca	agttgccaga	taatgtaagt	ctcgaggaag	120
gtgccatgtg	tgaaccactc	agtgttgggg	ttcatgcttg	tcgccgtgct	tctgtaggcc	180
ctgagacaaa	tgtcttggtg	atgggggacg	gtccatctcg	ccttgctacc	gtgctgtctg	240
cacgtgcatt	tggagcttca	cgaattatta	ttgctgatgt	agatgaagag	cgtctgtcaa	300
tggctaaaaa	ggttggctcc	gatgaatgag	tcttagtctc	cagagactct	caggatattg	360
atgaagaagt	gaccgcata	caaaatgccg	tgggtggaaa	catagatgta	acttttgatt	420
gtgctggttt	tgctaaaacc	atgtcgacgg	ctctaaagct	acgtctgctg	cggttaaggta	480
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&lt;210&gt; 348

&lt;211&gt; 352

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 348

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ctcactggaa	taaacactct	tcgcatccag	cccttcaaac	ttccgcctct	tgccccccat	120
gatgcgaagg	tgccgatgaa	ggctgtgggt	atctgtggca	gtgacgtcca	ctatttgagg	180
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ggaataattg	aggagggttg	cagtgaagtg	aaacatctgg	ttcctgggtg	ccgcgtagct	300
ttggagcctg	gaatatcgtg	ttggcggttg	gaccaatgta	agcgaggctc	ct	352

<210> 349  
 <211> 340  
 <212> DNA  
 <213> Pinus radiata

<400> 349  
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 catgaaggct gtgggtatct gtggcagtga cgtccactat ttgaggacat tacgggtgtgc 180  
 ggactttatt gtaaaagagc caatgggtgat tggcatgag tctgctggaa taattgagga 240  
 ggttggcagt gaagtgaac atctggttcc tggtagccgc gtagctttgg agcctggaat 300  
 atcgtgttgg cgttgtgacc aatgtaagcg aggtcctac 340

<210> 350  
 <211> 337  
 <212> DNA  
 <213> Pinus radiata

<400> 350  
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 gtgctcatga aggtgtggg tatctgtggc agtgacgtcc actatttgag gacattacgg 180  
 tgtgcggact ttattgtaaa agagccaatg gtgattggtc atgagctctg tggaataatt 240  
 gaggagggtg gcagtgaagt gaaacatctg gttcctggtg accgcgtagc tttggagcct 300  
 ggaatatcgt gttggcgttg tgaccaatgt aagcgag 337

<210> 351  
 <211> 500  
 <212> DNA  
 <213> Pinus radiata

<400> 351  
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 atgaaggctg tgggtatctg tggcagtga cgtccactat tgaggacatt acgggtgtgcg 180  
 gactttattg taaaagagcc aatgggtgatt ggtcatgagt ctgctggaat aattgaggag 240  
 gttggcagtg aagtgaacaa tctggttccg ggtgaccgcg tagctttgga gcctggaata 300  
 tcgtgttggc gttgtgacca atgtaagcga ggctcctaca atttgtgtcc cgagatgaag 360  
 ttttttgcaa cacctcccgt gcatgggttcc ttggccaatc agattgttca tcctgcagat 420  
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 agtgttgggg ttcattgcttg 500

<210> 352  
 <211> 589  
 <212> DNA  
 <213> Pinus radiata

<400> 352  
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 catgaaggct gtgggtatct gtggcagtga cgtccactat ttgaggacat tacgggtgtgc 180  
 ggactttatt gtaaaagagc caatgggtgat tggcatgag tctgctggaa taattgagga 240  
 ggttggcagt gaagtgaac atctggttcc tggtagccgc gtagctttgg agcctggaat 300  
 atcgtgttgg cggtgtgacc aatgtaagcg aggtcctac aatttgtgtc ccgagatgaa 360  
 gttttttgca acacctcccg tgcattggtt cttggccaat cagattgttc atcctgcaga 420  
 tttatgtttc aagttgccag ataatgtaag tctcgaggaa ggtgccatgt gtgaaccact 480

cagtgttggg gttcatgctt gtcgccgtgc ttctgtagge cctgagacaa atgtcttgg	540
aatgggggca ggtcctatcg gccttgtcac cgtgctgtct gcacgtgca	589

&lt;210&gt; 353

&lt;211&gt; 332

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 353

gacaatatgg ctgcatggct cactggaata aacactcttc gcatccagcc cttcaaactt	60
ccgctctctg gccccatga tgcgaagggt cgcgaagggt ctgtgggtat ctgtggcagt	120
gacgtccact atttgaggac attacgggtg gcggacttta ttgtaaaaga gccaatgggtg	180
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cctgggtgacc gcgtagcttt ggagcctgga atatcgtgtt ggcgttgtga ccaatgtaag	300
cgaggctcct acaatttgtg tcccagatg aa	332

&lt;210&gt; 354

&lt;211&gt; 312

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 354

gctcactgga ataaacactc ttcgcatcca gcccttcaaa ctccccctct tggcccccat	60
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acattacgggt gtgcggactt tattgtaaaa gagccaatgg tgatttggtca tgagtctgct	180
ggaataattg agggagggtg cagtgaagtg aaacatctgg ttcttgggtga ccgcgtagct	240
ttggagcctg gaatatcgtg ttggcggtgt gaccaatgta agcgaggctc ctacaatttg	300
tgtcccgaga tg	312

&lt;210&gt; 355

&lt;211&gt; 432

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 355

cttcccccg cttccaaggt agtgcagctg atcaagagcc aaggcatcaa caagttgaag	60
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ggcaatgaag tcttcgccgc ttctaattgac ctcacctcct acctgtctcc tgccatgaag	300
aatattcaca tggctctcgt caaatacaac ctcgacggaa ttatcaaggt gtcgagcccc	360
cttgccacca gtgtgctcca gaactctttc ccgccataac cggctttcaa gagcgacctt	420
gtggaaccac ga	432

&lt;210&gt; 356

&lt;211&gt; 384

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 356

taaggccact gaatgggtaa atgaaaacat tcgggcctac ttaccagcca caaagatcac	60
aggcatagct gtagggaacg aggtttacac aggaactgac acgcagttaa tggcaaacct	120
ggttcccgca atgaaaaaca tccattcggc ccttgctcagc atcgggtgcag acatgaatat	180
taaagttacc actccccatt ctcttgctgt acttgggaat tcatttccac cgtctgctgg	240
ttcctttgca tcaaactgta agagcctaata gaaaccactt ttggatttgt tgtctcagat	300
tggttctcct ttcttcataa atgcttatcc atattttgca tacaagggtg accccagcca	360
gatatccctg gcttatgtac tatt	384

<210> 357  
 <211> 420  
 <212> DNA  
 <213> Pinus radiata

<400> 357  
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 gatggtgata aaataggagt ggactatggc atggacgcaa gccatcttcc atctgcagac 120  
 gaggtggtaa ctttgatgaa gtccaacaac attgggaaaa ctagaattta ccaggaaaac 180  
 gatgtgttac tgcaagcttt cgcgaattct ggtatcgatg taatagtggg tgtcgctaac 240  
 gaagaactga agaacatata ttccagccaa gactncgcaa accgttgggt tagcgagcac 300  
 attgtgccct tctatcccgc caccaatgtc aaatacattg ctgtgggaaa cgagggtttg 360  
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<210> 358  
 <211> 399  
 <212> DNA  
 <213> Pinus radiata

<400> 358  
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 caacatccca ctcatcgtaa ctgaaagcgg atggccttct ggtggcaatg atgtggccac 180  
 ggttgacaac gctcgcgttt ataacaacaa tctcatccgc catgtgctct caaatgtagg 240  
 gactcccaag aggcgggaa cgagcattga gacctacatc ttgcgacttt tcaacgagaa 300  
 cagaaaagct ggtgatgaga cggagcgtca ctttgggctt ttctacctta accaacaatc 360  
 tgtatactct ctaaaacttta ctccgtaact gcgtcgcag 399

<210> 359  
 <211> 469  
 <212> DNA  
 <213> Pinus radiata

<400> 359  
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 caacatccca ctcatcgtaa ctgaaagcgg atggccttct ggtggcaatg atgtggccac 180  
 ggttgacaac gctcgcgttt ataacaacaa tctcatccgc catgtgctct caaatgtagg 240  
 gactcccaag aggcgggaa cgagcattga gacctacatc ttgcgacttt tcaacgagaa 300  
 cagaaaagct ggtgatgaga cggagcgtca ctttgggctt ttctacctta accaacaatc 360  
 tgtatactct ctaaaacttta ctccgtaact gcgtcgcagt ccgacgaacg aatagagcca 420  
 atatgaatat gtccctctata tgtcaactgc ctcgatagat atattatgt 469

<210> 360  
 <211> 473  
 <212> DNA  
 <213> Pinus radiata

<400> 360  
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 ctgttgctctg accatttttaa tgtacgttat ggcttcgttc caacgccagc cttgtacatt 120  
 gcaggccttt gcattctgca ttattgttct ttgctctttc tatgcagacg ctggaacagt 180  
 agggatttgc tatggacggg tggctgacaa tttggcggca ccggccgatg tggtaggcct 240  
 gctgaaaagac aacaacatca gcaaagtgcg gctcttcgac tcagaccctg cgggtgcttca 300  
 ggcttctgct gggctcgaga tcggactcat gacagctgtc cccaacgagt tggtaggag 360  
 catcggtagc aaccggagg ctgccgcagg gtgggtgcag ggaaacgttg tgcccttcca 420



cccggcgacc cggatcgaat acatcgcggt gggcaacgag gttttgcaca gca 473

<210> 361

<211> 441

<212> DNA

<213> Pinus radiata

<400> 361

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gccgaccgc	cacgggggac	acccggcgcc	cccggngcct	tcatccccac	ctatatatttc	420
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<210> 362

<211> 351

<212> DNA

<213> Pinus radiata

<400> 362

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gcaacccaag	caacattcat	ttggactacg	ctctctttca	gccacggct	acgccggtga	180
cagacaaaga	tcacagctac	agcaacctgt	tcgatgccat	ggttgatact	cttttgcgg	240
ccatggaagc	ctcgggggtat	cccaacatcc	cgatcgctcat	taccgaaagt	ggatggcctt	300
ctgctggcgc	ggaagtggcc	accattgaga	atgctcagac	ctataacaat	a	351

<210> 363

<211> 388

<212> DNA

<213> Pinus radiata

<400> 363

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gaacggccac	aacatcgga	aaatgaagct	tttcaatcca	gacgggtggg	cattgaatgc	180
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ctcctccagc	caggactcag	caaatgggtg	ggttaatgac	aatattgtgc	gctattccag	300
caccagtatc	aaatatattg	cggtgggcaa	cgagggtttt	cctagcacgc	agtaccgtat	360
cgtatcttgt	tccagccatg	aacaacat				388

<210> 364

<211> 560

<212> DNA

<213> Pinus radiata

<400> 364

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agtgggtctt	ttgcagacca	gcaatatcgg	cagaataaaa	ctctacactg	taaattgcgac	180
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gacttcatat	ccaccatctg	ctggttcatt	tcaaccagga	ctggagagcc	tactggaaca	540
gcttttggct	ctcctgtctc					560

&lt;210&gt; 365

&lt;211&gt; 494

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 365

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ttctccacc	atctgcagta	gcaaagctgg	ttcagagta	aagtatttca	aagttgagac	120
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attctctggc	catactttct	acatctgtcc	ccccatctgc	tggccgtttc	aatgaaagtt	480
ttgacatgaa	atcc					494

&lt;210&gt; 366

&lt;211&gt; 365

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 366

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aaggtaatag	ttgatctcca	tgtctgtcca	gggtcgcaaa	atgccgaatc	ttatagtgcg	360
acgag						365

&lt;210&gt; 367

&lt;211&gt; 435

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 367

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agaagcacia	tatcaagggt	atcgtcgacc	tgcatgggtg	cccaggcagt	cagaacgggt	180
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tgactgcaat	tgtcctccta	aatgagccag	caggatatct	tagccacgt	ctcctcgata	360
ctgcgaacaa	cattggctcg	nagttatggg	agtatccgca	ctccatttgg	caattcaacg	420
caaagccaca	tgctt					435

&lt;210&gt; 368

&lt;211&gt; 630

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 368

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tccagttgga	tggtggattg	ctagtgatcc	atatactcct	gtccttttgg	ttggaggatc	120

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catcaaaacc	acaagagctc	aaactctgca				630

&lt;210&gt; 369

&lt;211&gt; 507

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 369

aacggccttg	gacctgacaa	agctcctcaa	gtcatgaatg	atcactggaa	cagcttcatt	60
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&lt;210&gt; 370

&lt;211&gt; 480

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 370

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tgatccatat	cctcctgctc	cttttggttg	aggatctcta	gcttgctctg	acaaagcttt	360
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&lt;210&gt; 371

&lt;211&gt; 366

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 371

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gctgct						366

&lt;210&gt; 372

&lt;211&gt; 427

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 372

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tgatagcctt	ccggtacgnt	tgtgcctgag	cggtttgaag	cgagttaaac	tatccctgcc	360
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gagtgtg						427

&lt;210&gt; 373

&lt;211&gt; 384

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 373

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tgagaaaagt	aagggtcctt	aatcagtagt	tccgattaca	gatgaagagg	ttgttcttgg	300
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caactatggt	tctacgtata	cata				384

&lt;210&gt; 374

&lt;211&gt; 368

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 374

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cttaaaaaaa	tctcgtagct	ctttgtttgg	agcttagctc	agtggagtag	acttgcattc	360
aggcgggc						368

&lt;210&gt; 375

&lt;211&gt; 161

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 375

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&lt;210&gt; 376

&lt;211&gt; 283

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 376

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&lt;210&gt; 377

&lt;211&gt; 363

&lt;212&gt; DNA

<213> *Eucalyptus grandis*

&lt;400&gt; 377

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ttc						363

&lt;210&gt; 378

&lt;211&gt; 457

&lt;212&gt; DNA

<213> *Pinus radiata*

&lt;400&gt; 378

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tggtggcagt	tagtactaac	cttaagcttg	taaaagaatt	tggaaatagac	ccacaaaatg	360
cttttgcatt	ctgggattgg	gttggcggtc	gctacagcgt	gtgcagtgtg	gtgggtgccc	420
tccccttata	acttcagtat	gggtttccta	ttgttag			457

&lt;210&gt; 379

&lt;211&gt; 386

&lt;212&gt; DNA

<213> *Pinus radiata*

&lt;400&gt; 379

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tttgagaaaa	atattcctgt	tttgct				386

&lt;210&gt; 380

&lt;211&gt; 365

&lt;212&gt; DNA

<213> *Pinus radiata*

&lt;400&gt; 380

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ggtgaaatag	attttggaga	acctggtaca	aatggccagc	atagctttta	ccagttaatt	360
catca						365

&lt;210&gt; 381

&lt;211&gt; 491

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 381

tcagtgctg	tggtgcccct	ccccttatca	cttcaatatg	ggtttcctat	tgtagcaag	60
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catcagggtc	gtgtaattcc	ttgtgatttt	attggcatcg	tcaagagcca	gcagccatt	420
tatttacaag	gggaagttgt	gagtaaccac	gatgagctta	tgtctaactt	ttttgcacag	480
cggatgctc	t					491

&lt;210&gt; 382

&lt;211&gt; 446

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 382

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actgagtctt	gccaccatat	gcttcc				446

&lt;210&gt; 383

&lt;211&gt; 464

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 383

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&lt;210&gt; 384

&lt;211&gt; 385

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 384

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&lt;210&gt; 385

&lt;211&gt; 433

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 385

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&lt;210&gt; 386

&lt;211&gt; 349

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 386

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&lt;210&gt; 387

&lt;211&gt; 438

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 387

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&lt;210&gt; 388

&lt;211&gt; 414

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 388

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&lt;210&gt; 389

&lt;211&gt; 625

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 389

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&lt;210&gt; 390

&lt;211&gt; 78

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 390

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&lt;210&gt; 391

&lt;211&gt; 178

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 391

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&lt;210&gt; 392

&lt;211&gt; 359

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 392

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<210> 393

<211> 510

<212> DNA

<213> Eucalyptus grandis

<400> 393

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<210> 394

<211> 469

<212> DNA

<213> Pinus radiata

<400> 394

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<211> 443

<212> DNA

<213> Pinus radiata

<400> 395

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gcggtcaacg	gcggaatcga	caaagccaac	gtgttcttct	accacgctt	tggtctcttc	300
cggcagggcg	ttcatcacat	ggcctttcag	ctgcgtggta	ccgcgtacca	tgttttcaca	360
ggcgatgatg	ttcagcgggg	attcattacc	ttgttcttta	cgtttcacct	gccctttggc	420
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<210> 396

<211> 252

<212> DNA

<213> Pinus radiata

<400> 396

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cggttaactc	catgcctggg	atgttcggca	gtgcgccttt	gaactgcgtt	ttatcgacaa	180

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<210> 397

<211> 395

<212> DNA

<213> Pinus radiata

<400> 397

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catgcctggg atgttcggca gtgcgccttt gaactgcgtt ttatcgacaa tccattcgct 180  
gaagggtttct accgtcactt ccagcggatc gttagttgcc gaagccgaag gcggtacgat 240  
gcggtcaacg gcggaatcga caaagccaac gtgttcttct acccagcgtt tggcgtcttc 300  
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<210> 398

<211> 422

<212> DNA

<213> Pinus radiata

<400> 398

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<210> 399

<211> 305

<212> DNA

<213> Eucalyptus grandis

<400> 399

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caagggggccc ggtgcgaaca tcaacgccc agtgaagtgg cccggttaca agaagatcaa 240  
taagcaggaa gcagccaagt tcaccgctgg gacttttctg gatggggatt ggatcaaggg 300  
agcct 305

<210> 400

<211> 372

<212> DNA

<213> Eucalyptus grandis

<400> 400

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aagcaccagg ccgtggcgct gaggggtccag tcggacttct ccgtcttcta caactgccac 180  
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aaggagaggc gc 372

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 <212> DNA  
 <213> Eucalyptus grandis

<400> 401  
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 cacggcccag aaccgcaagg acccgaacca gaacacgggc atctcgattc acgcgtgccc 180  
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<210> 402  
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 <212> DNA  
 <213> Eucalyptus grandis

<400> 402  
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 caaagagaaa gtactggtag acaagaagac gaccaatgtc atgattatcg gcgacggccc 180  
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 <212> DNA  
 <213> Eucalyptus grandis

<400> 403  
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 ggagtggaac ggagactttg cgctagacac tttgtattac ggagagtaca tgaacgatgg 240  
 gcccggggca gccgtcgccc tacgtgtgaa atggccgggt ttccgagtca tcacatccac 300  
 aacagaggca aacaaattca cagtcgcgca gttcatattt ggatcttcat ggttgccgtc 360  
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 <213> Eucalyptus grandis

<400> 404  
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 acgattgcgg aggcgataaa gaaggcgccg gagagcagtg gccggaggat catcatctac 180  
 gtgcgagccg ggaggtacga ggaggataac ttgaagggtgg ggaagaagaa gacgaacctc 240  
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<210> 405

<211> 227  
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 ggccccaagg tcaacaggct gtcgccctaa ccgtggcctc cgatcacgcg gcattctata 180  
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<210> 406  
 <211> 373  
 <212> DNA  
 <213> Eucalyptus grandis

<400> 406  
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<210> 407  
 <211> 190  
 <212> DNA  
 <213> Eucalyptus grandis

<400> 407  
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 gtttgttaca tgttgtccta catgggcgat cacattcacc ccgaagggtg gctggagtgg 180  
 aacggagact 190

<210> 408  
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 <212> DNA  
 <213> Eucalyptus grandis

<400> 408  
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 gcaaacaat tcacagtcgc gcagttcata tttggatctt catggttgcc gtccaccggg 360  
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<210> 409  
 <211> 482  
 <212> DNA  
 <213> Eucalyptus grandis

<400> 409  
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caccgtcggga	gtgctcggcg	acggattcat	ggcaagcggg	ctcacgatcc	agaacaccgc	360
agggtccggtc	accaccagg	cgggtggcggt	ccggtcggac	agcgatttct	cggtcatcga	420
gaactgcgag	ttcttgggga	accaagacac	gctctacgcc	cactccctcc	ggcagtacta	480
ca						482

&lt;210&gt; 410

&lt;211&gt; 424

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 410

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tggt						424

&lt;210&gt; 411

&lt;211&gt; 519

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 411

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&lt;210&gt; 412

&lt;211&gt; 395

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 412

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acaccgcctc	cttcgcggcg	agcggagccg	gtttcattgc	ccgcgacatg	accttcgaga	360
actacgccgg	gccgggagaag	caccaggcgg	tggtc			395

&lt;210&gt; 413

&lt;211&gt; 499

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 413

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cattcaattc	cgcaactggt					499

&lt;210&gt; 414

&lt;211&gt; 497

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 414

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&lt;210&gt; 415

&lt;211&gt; 295

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 415

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gtttcattgc	ccgcgacatg	accttcgaga	actacgccgg	gccggagaag	cacca	295

&lt;210&gt; 416

&lt;211&gt; 433

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 416

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&lt;210&gt; 417

&lt;211&gt; 414

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 417

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&lt;210&gt; 418

&lt;211&gt; 382

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 418

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cgttccacac	cgcctccttc	tc				382

&lt;210&gt; 419

&lt;211&gt; 247

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 419

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tcaaccc						247

&lt;210&gt; 420

&lt;211&gt; 471

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 420

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&lt;210&gt; 421

&lt;211&gt; 371

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 421

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ggaagcccat g	371

&lt;210&gt; 422

&lt;211&gt; 349

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 422

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&lt;210&gt; 423

&lt;211&gt; 357

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 423

ggagggttgc gcaaatgccg gtgacggcga tacaggcgga cgtgacggta tcgaaggacg	60
ggaacgggac gtgcaagacg atctcggagg ccatcaagaa ggcgcgggac tacggtaccc	120
gccggtttat catatacgtg cgagccggaa ggtacgagga agataatctg aaggtgggga	180
ggaagaagac gaacgtgatg ttcgtagggg acgggaagag caacaccatc atctccggcg	240
gcaagagcat cttcgacaac atgacgacgt tccacaccgc gtccttcgct gccaccggag	300
ccgggttcat cgtcggggac atccgttcga gaactgggct gggcccgcca agcacca	357

&lt;210&gt; 424

&lt;211&gt; 346

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 424

gggggcggcg gcggcgagta tccagagtgg ctaggggaga gagagaggga gctgctggac	60
atgccggcgg cggaggtaca ggcggacata gtggtggcga aggacggagc gaacgggacg	120
tacaagacga tcgcggaggc gataaagaag gcgccggaga gcagtggccg gaggatcatc	180
atctacgtgc aagccgggag gtacgaagag gataaactga aggtggggaa gaagaagacg	240
aacctcatgt tcatcggcga tgggaagggc aaaacgggtca taacggggcg caaaagtgt	300
gccgacaaga tgaccacgtt ccacaccgcc tccttcgcgg cgagcg	346

&lt;210&gt; 425

&lt;211&gt; 577

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 425

gcgacggcgg gaccaccacc atcatcaccg gcgaccggac gtgaagggcg gcttcaccac	60
cttcgagtc gccaccgtcg cgggtggttg cgagcgattc ttggccaaaa gcataacctt	120



ccagaacacc	gntggccctt	caaaccacca	ggccgttgcg	ctccgggttg	gcgccgatct	180
atcagccttt	tacgaatgcg	acatcctcgc	ctaccaagat	accctctatg	tccacaacaa	240
ccgccaatte	tttgtcaagt	gcttaattgc	cggcacagtc	gacttcatct	ttggtaacgc	300
agctgtcgtc	atccaagact	gtgacatcca	tgcccgaag	ccaaaccctg	gccaaaagaa	360
catggttact	gctcaaggac	gaattgaccc	gaaccaaacc	acgggaatcg	tgatccaaaa	420
atgcaggatt	gtgagacca	acgatctccg	atcagtgaag	agcagtttcc	caacgtacct	480
cggtcgtcca	tggaaggagt	actcgaggac	agtgattatg	caatcatcga	tctcggacgt	540
aatcgaccgc	gtgggttggc	acgagtggag	tgggacc			577

&lt;210&gt; 426

&lt;211&gt; 283

&lt;212&gt; DNA

<213> *Eucalyptus grandis*

&lt;400&gt; 426

aaagaccggg	gtgtacagag	agaacgtgca	ggtgccgaag	aagaagacca	acctgacttt	60
catcggcgac	ggcgaggacca	ccaccatcat	caccggcgac	cggagcgtga	agggcggttt	120
caccaccttc	gagtccgccca	ctgtcgcggg	gcttggcgag	cgattcttgg	ccaaaaacat	180
aaccttccag	aacaccgcgc	gcccttcaaa	ccaccaggcc	gttgccctcc	gtgttggtgc	240
cgatctatct	gccatttacg	aatgcgacat	cctcgcctac	cag		283

&lt;210&gt; 427

&lt;211&gt; 345

&lt;212&gt; DNA

<213> *Eucalyptus grandis*

&lt;400&gt; 427

ccaccgtcgg	agtgccttggc	gacggattca	tggcaaccgg	gctcacgata	cagaacaccg	60
cgggtccaga	cgcccaccag	gcggtggcat	tccggtcgga	cagcgatttc	tcggtcacgc	120
agaactgcga	gttcttgga	aaccaggaca	cgctctatgc	ccacgccttc	cggcagtact	180
acaagtccctg	ccacatcgag	ggcaatgtgg	acttcatctt	tgggaactcg	gcctcctact	240
tccaggactg	ccagatccctg	gtccgcccc	ggcagggtcaa	gcccagagaag	ggcgagagca	300
atgctgtcac	agcccatggc	cggaccgacc	ccgcgcagtc	gacag		345

&lt;210&gt; 428

&lt;211&gt; 478

&lt;212&gt; DNA

<213> *Eucalyptus grandis*

&lt;400&gt; 428

tgagcagttc	actcacaat	tacatcacct	gcttggacgg	tttcgaaggc	tcgtcatcag	60
ctaaatcttc	gatcaagcct	attctcagcg	acttgatata	gagggcaaga	acttctctag	120
ccatatttgt	ttctacttca	tctcctgaag	gacgaagacc	agatgttctg	gagtccttga	180
tcggtgattt	cccacatgg	gtcacacgaa	aagatcatcg	tctcctgcaa	tctctggtga	240
acgcagttaa	tgccgacgtg	gtggtggcga	aggacggaac	tgggaagtgc	aagacagtga	300
aagaggcgat	cgcagctgct	cctagcaaa	cccagacccg	gtacgttatt	tatgtgaaga	360
aaggcacata	caaggagaat	gtggagggtg	caaagacaaa	gacaaacatc	atgcttggtg	420
gcgacggcat	ggattcaact	gtgatcactg	gcagcctcaa	cgtcattgac	ggtgcgac	478

&lt;210&gt; 429

&lt;211&gt; 335

&lt;212&gt; DNA

<213> *Eucalyptus grandis*

&lt;400&gt; 429

tttttttttt	tttttttttt	taatcnccaa	acattocaaa	ttattgaatt	cacaacacaa	60
cttacagaaa	ctccccaga	aagaaaggac	caatactcat	ttttcacagc	ccttcagtg	120

aagacactcc	agtggacttc	aaccaagtcc	ctccctgaat	cagctcggcc	acggngaact	180
tcttcgcctc	tgctgggctg	gtgatgacat	gataacctgg	ccacttcacc	cgcttgctcg	240
ttcccgcgcc	tggtcccttg	ttcatgtact	ccccgtagta	canagtcttg	agggcatgat	300
cgccgctcca	caccgaccac	cctgtgggat	caatg			335

&lt;210&gt; 430

&lt;211&gt; 361

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 430

tttttttttt	tttttttttt	taatctccaa	acattccaaa	ttattgaatt	cacaacacaa	60
cttacagaaa	ctccccaga	aagaaaggac	caatactcat	ttttcacagc	ccttcagtgt	120
aagacactcc	agtggacttc	aaccaagtcc	ctccctgaat	cagctcggcc	acggngaact	180
tcttcgcctc	tgctgggctg	gtgatgacat	gataacctgg	ccacttcacc	cgcttgctcg	240
ttcccgcgcc	tggtcccttg	ttcatgtact	ccccgtagta	cagagtcttg	agggcatgat	300
cgccgctcca	caccgaccac	cctgtgggat	caatgtgatc	accaatgttt	gattgcatca	360
c						361

&lt;210&gt; 431

&lt;211&gt; 368

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 431

tttttttttt	ttttttacga	tcaaattctcc	aaacattcca	aattattgaa	ttcacaacac	60
aacttacaga	aactccccca	gaaagaaagg	accaatactc	atttttcaca	gcccttcagt	120
gtaagacact	ccagtggact	tcaaccaagt	ccctccctga	atcagctcgg	ccacggtgaa	180
cttcttcgcc	tctgctgggc	tggtgatgac	atgataacct	ggccacttca	cccgttgct	240
cgttcccgcg	cctgggtccct	tggtcatgta	ctccccgtag	tacagagtct	tgagggcatg	300
atcgccgctc	cacaccgacc	accctgtggg	atcaatgtga	tcaccaatgt	ttgattgcat	360
ccacagct						368

&lt;210&gt; 432

&lt;211&gt; 324

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 432

cccaaccaga	acaccggcat	ttcgatccat	gcttgccaga	tcgtcgccgc	tccagatctc	60
gaggcatcta	aaggaagcat	cccgcgtac	ctcgggcggc	catggaagat	gtactcgagg	120
gttgtgtaca	tggtgtccta	catgggcgat	cacattcacc	ccgaagggtg	gctggagtgg	180
aacgggagact	ttgcgctaga	cactttgtat	tacggagagt	acatgaacga	tgggcctggg	240
gcagccgctg	gcctacgtgt	gaaatggcct	ggtttccgag	tcatcacatc	cacaacagag	300
gcaaacaaat	tcacagtgcg	gcag				324

&lt;210&gt; 433

&lt;211&gt; 460

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 433

gcccgcgggc	ggcgggggcg	agtatccaga	gtggctaggg	gagagagaga	gggagctgct	60
ggacatgccg	gcgggcgagg	tacaggcgga	catagtgggtg	gcgaaggacg	gagcgagcgg	120
gacgtacaag	acgattgcgg	aggcgataaa	gaaggcgccg	gagagcagtg	gccggaggat	180
catcatctac	gtgcgagccg	ggaggtacga	ggaggataac	ttgaagggtg	ggaagaagaa	240
gacgaacctc	atgttcatcg	gcgatgggaa	gggcagaaacg	gtcataaacg	gcggcaaaaa	300

tgtagccgac	aagatgacca	cgttccacac	cgctccttc	ggtaaatttc	tgtgtccata	360
tccgaatttc	taatgttcaa	actctcgact	aagctaggcc	aaaaattata	aataatcttt	420
tttgtctaaa	taatttattt	tttacgaaac	aaatcgaacc			460

&lt;210&gt; 434

&lt;211&gt; 344

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 434

aggacggaac	tgggaagtgc	aagacagtga	aagaggcgat	cgcagctgct	cctagcaaag	60
cccagaccg	gtacgttatt	tatgtgaaga	aaggcacata	caaggagaat	gtggagggtg	120
caaagacaaa	gacaaacatc	atgcttggtg	gcgacggcat	ggattcaact	gtgatcactg	180
gcagcctcaa	cgtcattgac	ggcgcgacaa	cattcaattc	cgcaactgtt	gctgtgaatg	240
gcgatgggtt	catagcccag	gacatatggt	tccagaacac	tgccgggccg	cagaaacacc	300
aggccgctgc	actccgtgtc	agtgcagaca	agtcagtcac	caac		344

&lt;210&gt; 435

&lt;211&gt; 295

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 435

acgagctcga	gacatgacct	tcgagaacac	ggaaggaccc	gcgaacacca	ggcgggtggcc	60
ctgcgtgtgg	gatcagatct	ctcggctttc	tatcgctgca	gcttcaaggg	ttaccaggac	120
accctttacg	ccattccct	tcgtcagttt	tacagagaat	gcaacatcta	tggcaccgta	180
gatttcacat	tcggcaactc	cgccgctcgt	tttcaggatt	gcaatttgct	ggcgcggaga	240
cccctggaga	atcagacgat	tctttacacc	gctcagggca	ggcaggaccc	caatg	295

&lt;210&gt; 436

&lt;211&gt; 332

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 436

tgcaggtatt	tgttgacagt	gacatgagct	ttatgaacag	tgcagggcct	gacaagcatc	60
aagctgtggc	tctacgggtg	ggggccgatt	ttgcagcgat	ttatcgatgc	agtattattg	120
gttaccaaga	cacactttat	gttcaactct	tgaggcagtt	ttacagagaa	tgtgacgtgt	180
tcggaacagt	ggacttcatt	tttggcaatg	cagccgtggt	tttacaggag	tgtaacattt	240
atgctcgaca	aggcatgccc	aatcaagtga	atgtaatcac	tgcccaagga	aggaatcatc	300
cttatcaaaa	taccggcatc	tcaatacata	at			332

&lt;210&gt; 437

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 437

gcggcccccag	agaaaagtgg	taagagatat	gtgatcaagg	tgaagaaggg	aacgtttacaa	60
ggagaacgtg	gaggtgggtg	aaacgaagac	taatcatcatg	ttgattggag	aaggcatgga	120
ggccacaatc	gttacaggga	gcagaaatgt	gatagacgga	tccaccactt	tcaattcagc	180
cacattcgct	gctgtagggg	agggatttat	ggcacaagac	atggcgttcg	tcaacacagc	240
aggcccgagc	aaacatcagg	cgggtggctct	tcgagtaggt	cagatcaatc	agtgttatat	300
c						301

&lt;210&gt; 438

&lt;211&gt; 242

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 438

gaagaaggga	acgtacaagg	agaacgtgga	ggtgggtaaa	acgaagacta	atatcatgtt	60
gattggagaa	ggcatggagg	ccacaatcgt	tacagggagc	agaaatgtga	tagacggatc	120
caccactttc	aattcagcca	cattcgctgc	tgtagggaag	ggatttatgg	cacaagacat	180
ggcgcttcgtc	aacacagcag	gcccggacaa	acatcaggcg	gtggctcttc	gagtaggatac	240
ag						242

&lt;210&gt; 439

&lt;211&gt; 255

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 439

acttaactcc	ggtggaagg	aaccatgaat	ctccttcaat	gaattttgag	actgtaaagt	60
aactggcctc	gcttgagcta	ttaatcactc	gataccccgc	ccatttgacc	cgattttag	120
tgcccgcacc	aggccgcgg	ttcatgtatt	ccccatagta	taggtgctg	agagcgaagg	180
atccattcca	ttcgagccaa	ccggcagggt	gaatcaggtc	gccagaaaag	gactgcatga	240
agacagtgcg	agagt					255

&lt;210&gt; 440

&lt;211&gt; 362

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 440

gtggactgca	gccatgaatc	tccttgatg	aattctccaa	cagtaaactt	gctcgctct	60
tgagaacttg	tgattaccgc	ataccggcc	catttcacgc	ggtagcagt	tgctgctct	120
gggcctgtat	tcattgtattc	tccataatac	aaagtgtgca	atgcaaaact	tccattccat	180
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tcggaatcgg	gagtgatcct	gcagttgtga	atggaagtcc	ctgtgttctg	gttcggatca	360
gt						362

&lt;210&gt; 441

&lt;211&gt; 286

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 441

aagaaaacta	atatcatgtt	cgttggagat	ggtatggatg	tcacagtgg	gaccggaaac	60
cgaaatgtga	aggacaattt	cacaaccttt	cgttctgcaa	ctgttgctgt	gactggaaat	120
ggattcatcg	ctcgcgacat	gaccttcgag	aacacggcag	gaccgcgcaa	gcaccaggcg	180
gtggccctgc	gtgtgggatac	agatctttcg	gctttctatc	gatgcagctt	caagggttac	240
caggacaccc	tttacgceca	ttcccttcgt	caagttttta	cagaga		286

&lt;210&gt; 442

&lt;211&gt; 302

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 442

ggagaagagc	cagacgagat	acgtaattca	tataaaagca	ggagtattatg	cagagaatgt	60
ggagtgtcac	aagaagaaaa	ctaatatcat	gttcgttgga	gatggtatgg	atgtcacagt	120
ggtgaccgga	aaccgaaatg	tgaaggacaa	tttcacaacc	tttcgttctg	caactgttgc	180

tgtgactgga aatggattca tcgctcgcca catgaccttc gagaacacgg caggacccgc	240
gaagcaccag gcggtggccc tgcgtgtggg atcagatctt tcggctttct atcgatgcag	300
ct	302

<210> 443  
 <211> 466  
 <212> DNA  
 <213> Pinus radiata

<400> 443	
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cgaactatcg tcaccggatc attgtctgcc cagattcccg gcgtcggcac acacggctct	120
gcaactgtcg ggggtgaacgc ggacggtttt gtgcgcgag acattgcgtt cgagaatact	180
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gagaggtgag cctttctcgg acaccaggac accctatacg cgcacgccct ccgccagttt	300
tatcggaatt gcaggatcga aggcaacgtc gacttcatat ttggaaacgc agcggccatc	360
ttccacaact gctccattct cgtccgccct cgcaggtgc cgtctaactt ttccgaagcg	420
aaccccataa ctgcccacgg gcgattggat ccgggtcaga ctactg	466

<210> 444  
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 <212> DNA  
 <213> Pinus radiata

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aaaaattgaa agcgaactac agacacagac gaacttcttc aaaagtttgg cgaagtcgct	120
aatttatcaa tccgtcgatg tagtcgatgc cagtggactg caaccaggaa tttccttgta	180
tgaattctcc cactgtaaatt ttgctgcct cttgagaact tttgatcacc cgatacccag	240
gccatttcac acggtttcca gtagcggagc ctggccctgt attcatatat tctccataat	300
acaaagtttt caaagcaaaa gtcctatccc attccagcca gcccg	345

<210> 445  
 <211> 183  
 <212> DNA  
 <213> Pinus radiata

<400> 445	
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tatttgggaa tgccgcagtg gtgtttcaga gctgcaatct gatgcccgag aaacccgggtg	120
caaatcagaa aaatgccatc acagcacagg gcagaactga tccgaaccaa aacacaggaa	180
ctt	183

<210> 446  
 <211> 264  
 <212> DNA  
 <213> Pinus radiata

<400> 446	
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tcctacagc agcgaatgtg gctgaattga aagtgggtga tccgtctatc acatttctgc	180
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<210> 447  
 <211> 417

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 447

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atgggctgtat	agggtgtcct	ggtaaccctt	gaagctgcaa	cgatagaaag	ccgagagatc	120
tgatcccaca	cgcagggccca	cgcctgggtg	cttcgcgggt	cctgccgtgt	tctcgaagg	180
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gaacatgata	ttagttttct	tcttgtgcaa	ctccacattc	tctgcataaa	ctcctgcttt	360
tatatgaatt	acgtatctcg	tctggctctt	ctccggagcc	ttctccactg	cctctgt	417

&lt;210&gt; 448

&lt;211&gt; 404

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 448

attatatctc	agctgacata	aatatatatt	aaaaattaca	gcatacaatg	gaactttgac	60
tgacagcgag	agaaaagaaa	gcccttaatt	tattggccac	ctggcttgca	atattctatt	120
gcattttaga	gtacagcaga	ataatataca	cgtcagcctt	aatttcagaa	aaataaataa	180
actacgtgag	cagccctcca	ataaaaaaga	tcccagtggg	tggaaccat	ttagcaccag	240
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caggccacgt	caccgcctc	gcaagccctg	cgccggggcc	actgttcata	tattcccat	360
aaaataaggt	atccagtgcg	aagtctccga	accactccaa	ccat		404

&lt;210&gt; 449

&lt;211&gt; 173

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 449

gcctcgctgg	agctcttaat	cgctcgatat	cctggccatt	tgaccgggcc	tgcatgtttt	60
gcgccaagggc	cgcggttcat	gtattcccca	taataagagg	tactcagtgc	gaaggttcca	120
ttccattcga	gccaaaccgc	tggtgaatc	atgtcgccaa	gaaacgattg	cat	173

&lt;210&gt; 450

&lt;211&gt; 398

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 450

ggctctgaga	gtaggagcgg	atthttgcagc	cttttacgt	tgacaggtca	tcggttacca	60
ggacacactg	tacgtacatt	ctctccgccca	atthttacaga	gaatgcgaca	tctacggcac	120
agtggacttc	atctttggca	acgcagccgt	ggtgttgag	aagtgcacca	tggtcggcag	180
aaaacccctg	cccaactcca	agatcacggt	gacggctcag	ggcaggaagg	acccaacca	240
gaacaccggc	atctccatcc	acgactgcag	agtgcggcg	gcggcggatc	ttgctccgt	300
caagggcctc	tatcgcgctt	acctcgggag	gccttggaag	ttatactctc	gcacggtata	360
cctgcaaaact	tttttgatg	atattattga	ccctgccg			398

&lt;210&gt; 451

&lt;211&gt; 404

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 451

agagaatgca	acatctatgg	caccgtagat	gttgcatctt	ctgtaaaact	gacgaaggga	60
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atgggcgtat	aggggtgtcct	ggtaaccctt	gaagctgcaa	cgatagaaaag	ccgagagatc	120
tgatcccaca	cgcagggcca	ccgcctgggtg	cttcgcgggt	cctgccgtgt	tctcgaagggt	180
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gaaattgtcc	ttcacatttc	ggtttccggt	caccactgtg	acatccatac	catctccaac	300
gaacatgata	ttagttttct	tcttgtgcaa	ctccacattc	tctgcataaa	ctcctgcttt	360
tatatgaatt	acgtatctcg	tctggctctt	ctccggagcc	ttct		404

&lt;210&gt; 452

&lt;211&gt; 394

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 452

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gaacatgata	ttagttttct	tcttgtgcaa	ctccacattc	tctgcataaa	ctcctgcttt	360
tatatgaatt	acgtatctcg	tctggctctt	ctcc			394

&lt;210&gt; 453

&lt;211&gt; 428

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 453

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atcatcct						428

&lt;210&gt; 454

&lt;211&gt; 329

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 454

gcaaatacga	aaaatgcaat	caccgcacag	ggcagaactg	atccgaacca	gaacacagga	60
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caaactttgt	attatggaga	atacatgaat	acaggcccag	gagcagcaac	tgctaaccgc	300
gtgaaatggg	ccgggtatcg	ggtaatcac				329

&lt;210&gt; 455

&lt;211&gt; 358

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 455

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gtgatcaagg tgaagaagg aacgtacaag gagaacgtgg aggtgggtaa aacgaagact	180
aatatcatgt tgattggaga aggcattggag gccacaatcg ttacagggag cagaaatgtg	240
atagacggat ccaccacttt caattcagcc acattcgctg ctgtagggaa gggatttatg	300
gcacaagaca tggcgttcgt caacacagca ggcccggaca aacatcaggc ggtggctc	358

&lt;210&gt; 456

&lt;211&gt; 195

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 456

cgtggagggtg ggtaaaacga agactaatat catgttgatt ggagaaggca tggaggccac	60
aatcggttaca gggagcagaa atgtgataga cggatccacc actttcaatt cagccacatt	120
cgctgctgta ggggaaggat ttatggcaca agacatggcg ttcgtcaaca cagcaggccc	180
ggacaaacat caggc	195

&lt;210&gt; 457

&lt;211&gt; 405

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 457

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cgtgatcgtg gctaaggatg gctctggaaa tttcaaaacg atttcacaag ccatagctgc	240
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gaacgtggag gtgggtaaaa cgaagactaa tatcatgttg attggagaag gcatggaggc	360
cacaatcggt acaggggagca gaaatgtgat agacggatcc accac	405

&lt;210&gt; 458

&lt;211&gt; 326

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 458

cgtggaacat ggatggccta caagacactc gacagcaaac gaaacatagt caaaacatta	60
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attaccgatg acccggccca tttcacgagg ttagcagttg ctgctcctgg gcctgtattc	240
atgtattctc cataatacaa agtttgcaat gcaaaacttc cattccattc cagccagccc	300
gcaggctgaa taacatcgtc tagata	326

&lt;210&gt; 459

&lt;211&gt; 360

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 459

tcgacctacg atcctctcca aacaacgtag tgccaaacgt gatcgtggct aaggatggct	60
ctggaaatct caaaacgatt tcacaagcca tagctgcggc cccagagaaa agtggttaaga	120
gatatgtgat caaggatgaag aagggaacgt acaaggagaa cgtggagggtg ggtaaaacga	180
agactaatat catgttgatt ggagaaggca tggaggccac aatcggttaca gggagcagaa	240
tgtgatagac ggatcnccac tttcaattca gccacattcg ctgctgtagg gaagggattt	300
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&lt;210&gt; 460



<211> 438  
 <212> DNA  
 <213> Pinus radiata

<400> 460  
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 attacccgat acccgggcca ttccacgcgg ttagcagttg ctgctcctgg gcctgtattc 240  
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 gcaggctgaa taacatcgtc tagataggac tgcattgaaa ccgttcggga atactccttc 360  
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 gtgatcttgc agttgtga 438

<210> 461  
 <211> 380  
 <212> DNA  
 <213> Pinus radiata

<400> 461  
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 tgatcccaca cgcagggcca ccgcctggtg cttcgcgggt cctgccgtgt tctcgaagggt 180  
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 gaaattgtcc ttcacatttc ggtttcgggt caccactgtg acatccatac catctccaac 300  
 gaacatgata ttaagtttcc ttcttggtgca actccacatt ctctgcataa actcctgctt 360  
 tatatgaata cgtatctcgt 380

<210> 462  
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 <212> DNA  
 <213> Pinus radiata

<400> 462  
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 gaaattgtcc ttcacatttc ggtttcgggt caccactgtg acatccatac catctccaac 300  
 gaacatgata ttagtgttct tcttggtgcaa ctccacattc tctgcataaa ctctgcttt 360  
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<210> 463  
 <211> 441  
 <212> DNA  
 <213> Pinus radiata

<400> 463  
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 tataacagcg ctactgttgc agtcaattcg ccttacttca tcgccagaaa cattacgttt 180  
 canaacacag ccccggttcc tctgccgggg gcggtgggca gacaagcagt ggccttgaga 240  
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 tatgaccacg ctggccgcca ttatttcaaa gactgttata tcgaaggctc cgtcgatttc 360  
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 tacggcgcgg tgacagcgca g 441

<210> 464  
 <211> 481  
 <212> DNA  
 <213> Pinus radiata

<400> 464  
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 gggtttacgg ggaaacgggt cgaatcccc gcagtaagac gaatctcgtg tttgtgggcg 180  
 ccggtatgga taaaacgggt atcacccggt ctgcatatgt gccgtctctg cccggcccg 240  
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 taacattccg aaacacattt caggggccac agactcatca agccgtggcc ctgagagtag 360  
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 t 481

<210> 465  
 <211> 505  
 <212> DNA  
 <213> Pinus radiata

<400> 465  
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 cccttcaggg tcatgattat catgtacaga caaacaactt tacagtccca tcgtctctctc 120  
 caacaaaaag gcgacgcctt ttggcagagg caggggaaga aatgaacaat gctcttcgga 180  
 atcaagaatt ttatgaccat tatggattga ttcattggagg ggcgcagcat gaatttctc 240  
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 atgcaccttc cctgctaagc agcagaaggt atatcatcta tgtgaaaaca ggcgtgtata 420  
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 gaaaaactat tgtagcagca ggcaa 505

<210> 466  
 <211> 361  
 <212> DNA  
 <213> Pinus radiata

<400> 466  
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 aagagatacg tgatcaaggt gaagaagggg acgtacaagg agaacgtgga agtgggcaaa 180  
 aagaagacaa atattatgct gatcgganaa ggcatggaag ccacgatcgt tacagggagc 240  
 agaaatgttg tagacgggtc caccactttc aattcctcta cactagctgc ttaggggaag 300  
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 g 361

<210> 467  
 <211> 402  
 <212> DNA  
 <213> Pinus radiata

<400> 467  
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 ctccagatgg ctctggaaaa ttcaaaacaa ttacagaagc catagctgcg gccccggaga 180  
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tgaggcaaaaa	gaagacaaat	attatgctga	tcggagaagg	catggaagcc	acgatcggtta	300
cagggagcag	aaatgttgta	gacgggtcca	ccactttcaa	ttcctctaca	ctagctgctg	360
tagggaaggg	gtttatggca	caagacatgg	cggttcgtcaa	ca		402

&lt;210&gt; 468

&lt;211&gt; 397

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 468

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taacattccg	aaacacattt	cagggggccac	agactcatca	agccgtggcc	ctgagagtag	360
acagcgattt	ttctgccttc	tacagctgcg	ctttcga			397

&lt;210&gt; 469

&lt;211&gt; 349

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 469

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gctcacggca	ggcaggaccc	caatgagaac	actggaattt	ccattcagaa	ctgtaaatgtg	300
accgcagccc	cagacctggc	tccagtgaag	agctcgttcg	atgcatatc		349

&lt;210&gt; 470

&lt;211&gt; 375

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 470

cggaggctcc	tacgtctcct	tccaaacgac	atcgtggctg	acgtgatcgt	ggctcaggat	60
ggctctggaa	aattcaaaac	aattacagaa	gccatagctg	cggccccgga	gaaaagctct	120
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gtggctcttc	gtgta					375

&lt;210&gt; 471

&lt;211&gt; 367

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 471

tgatccctga	cattactgtg	tcaaagctgg	atcagaaatc	ctctctatca	agcattcagc	60
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367

<210> 472  
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 <212> DNA  
 <213> Pinus radiata

&lt;400&gt; 472

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<210> 473  
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 <212> DNA  
 <213> Pinus radiata

&lt;400&gt; 473

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ggctccgctg	atttcacctt	cgggaacggc	ctctccctct	acgag		345

<210> 474  
 <211> 268  
 <212> DNA  
 <213> Pinus radiata

&lt;400&gt; 474

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catcgctcgc	gacatgacct	tcgagaacac	ggcaggaccc	gcgaagcacc	aggcgggtggc	180
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caccctttac	gccattccc	ttcgtag				268

<210> 475  
 <211> 316  
 <212> DNA  
 <213> Pinus radiata

&lt;400&gt; 475

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<210> 476  
 <211> 440

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 476

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caacatctat	ggcaccatag	atttcattct	cggcaactcc	gccgtcgttt	ttcaggattg	420
caatctcctg	gcgcggaggc					440

&lt;210&gt; 477

&lt;211&gt; 357

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 477

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&lt;210&gt; 478

&lt;211&gt; 318

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 478

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gcggggtatc	aggtgattaa	gaaatccaag	gaggccaaga	aatttacagt	gtctcaattc	300
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&lt;210&gt; 479

&lt;211&gt; 271

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 479

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ngacattgtc	cgtcagcatt	ggngacata	tggtcgtcat	tattatacca	gatatgatta	240
tgagaatggt	gattcaggag	cagcaaagga	a			271

&lt;210&gt; 480

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 480

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&lt;210&gt; 481

&lt;211&gt; 287

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 481

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&lt;210&gt; 482

&lt;211&gt; 285

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 482

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&lt;210&gt; 483

&lt;211&gt; 427

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 483

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&lt;210&gt; 484

&lt;211&gt; 408

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 484

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<212> DNA  
<213> Eucalyptus grandis

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<212> DNA  
<213> Eucalyptus grandis

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<213> Eucalyptus grandis

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<212> DNA  
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<400> 492  
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&lt;210&gt; 493

&lt;211&gt; 424

&lt;212&gt; DNA

<213> *Eucalyptus grandis*

&lt;400&gt; 493

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&lt;210&gt; 494

&lt;211&gt; 257

&lt;212&gt; DNA

<213> *Eucalyptus grandis*

&lt;400&gt; 494

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&lt;210&gt; 495

&lt;211&gt; 483

&lt;212&gt; DNA

<213> *Eucalyptus grandis*

&lt;400&gt; 495

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&lt;210&gt; 496

&lt;211&gt; 353

&lt;212&gt; DNA

<213> *Eucalyptus grandis*

&lt;400&gt; 496

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&lt;210&gt; 497

&lt;211&gt; 442

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 497

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&lt;210&gt; 498

&lt;211&gt; 364

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 498

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catt						364

&lt;210&gt; 499

&lt;211&gt; 365

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 499

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catct						365

&lt;210&gt; 500

&lt;211&gt; 390

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 500

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 <212> DNA  
 <213> Eucalyptus grandis

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&lt;210&gt; 505

&lt;211&gt; 477

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 505

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&lt;210&gt; 506

&lt;211&gt; 436

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 506

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&lt;210&gt; 507

&lt;211&gt; 473

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 507

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&lt;210&gt; 508

&lt;211&gt; 379

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 508

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&lt;210&gt; 509

&lt;211&gt; 459

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 509

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&lt;210&gt; 510

&lt;211&gt; 268

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 510

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&lt;210&gt; 511

&lt;211&gt; 293

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 511

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&lt;210&gt; 512

&lt;211&gt; 423

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 512

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&lt;210&gt; 513

&lt;211&gt; 508

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 513

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&lt;210&gt; 514

&lt;211&gt; 502

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 514

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&lt;210&gt; 515

&lt;211&gt; 447

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 515

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&lt;210&gt; 516

&lt;211&gt; 403

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 516

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&lt;210&gt; 517

&lt;211&gt; 379

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 517

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&lt;210&gt; 518

&lt;211&gt; 404

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 518

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&lt;210&gt; 519

&lt;211&gt; 705

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 519

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&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 524

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&lt;210&gt; 525

&lt;211&gt; 363

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 525

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&lt;210&gt; 526

&lt;211&gt; 344

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 526

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&lt;211&gt; 445

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 527

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&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 528

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&lt;210&gt; 529

&lt;211&gt; 505

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 529

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&lt;210&gt; 530

&lt;211&gt; 540

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 530

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&lt;210&gt; 531

&lt;211&gt; 398

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 531

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ggagacatgg	attggtcaac	agaaggggaa	gaagctttac	cttgatttaa	taagtcttca	360
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 aacgtgaaag gggtcgcaga gaagcgactg cggacatgtc tgaggactta tctgagggag 180  
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 <213> Eucalyptus grandis

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 gaagcccaga gaaaagctaa gcgtcgactt gaacgtgaaa ggggtcgcag agaagcgact 180  
 gcggacatgt ctgaggactt atctgaggga gaaaaggggtg atgcagtcag cgatatatcg 240  
 actcatggtg atagcaacag aggcagattg cctaggataa gttctgttga tgcaatggag 300  
 acatggattg gtcaacagaa ggggaagaag ctttaccttg tattaataag tcttcattggc 360  
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 <212> DNA  
 <213> Eucalyptus grandis

<400> 534  
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 caatacgaca tacaagataa tgcgccgaat agaggcagaa gaactgtccc ttgattcctc 180  
 tgagatagtg ataactagta ctaggcagga gattgatgag caatggcgct tgtatgatgc 240  
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 <212> DNA  
 <213> Eucalyptus grandis

<400> 535  
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 cgatcctcga cgtcggcccc gcgctcgacg acaagaaatc gtcgctgctg ctgcgggagc 180  
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<210> 536  
 <211> 441  
 <212> DNA  
 <213> Eucalyptus grandis

<400> 536  
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cgccataact	ccaatcaaca	tctgggtgatg	atatctgtct	agtaagcaag	tccactcgat	420
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&lt;210&gt; 537

&lt;211&gt; 389

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 537

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ttcagcaatg	tgaaagtaga	agattcaact	gaaagcgaga	gcgacctcaa	atccttgatt	240
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aatgtcacaa	ccttgctgaa	ggcttttcgg				389

&lt;210&gt; 538

&lt;211&gt; 647

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 538

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&lt;210&gt; 539

&lt;211&gt; 340

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 539

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ctctaccgct	cctgggtcaa	ggcccaggcg	acgaggagcc	cgcagganag	gaacacgcga	300
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&lt;210&gt; 540

&lt;211&gt; 320

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

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 cggaggagac gtctcgttcg 320

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 <212> DNA  
 <213> *Eucalyptus grandis*

<400> 541  
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 gagcaagttg gcggtggcca accagtgtgg cctgttgcca tccatggtea ctatgctgat 180  
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 cactcacttg gtcgagataa gttggaacaa ctctcaagc aaggccgtct gtccagggat 300  
 gaaatcaata cgacatacaa gataatgcgc cgaatagagg cagaagaact gtcccttgat 360  
 tctctgagat agtgataact atacta 386

<210> 542  
 <211> 326  
 <212> DNA  
 <213> *Eucalyptus grandis*

<400> 542  
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<210> 543  
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 <212> DNA  
 <213> *Eucalyptus grandis*

<400> 543  
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 <211> 558  
 <212> DNA  
 <213> *Eucalyptus grandis*

<400> 544  
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caatggagac	atgggttg					558

&lt;210&gt; 545

&lt;211&gt; 414

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 545

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&lt;210&gt; 546

&lt;211&gt; 289

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 546

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cgcgcccgct	tcagcccacc	cgctacttcg	tcgaggaggt	catcaccggc	ttcgatgaga	240
ccgatctcta	ccgctcctgg	gtcaaggccc	aggcgacgag	gagcccga		289

&lt;210&gt; 547

&lt;211&gt; 227

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 547

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cgggggtgaaa	acatggagct	tggccgagac	tcggatactg	gtggtca		227

&lt;210&gt; 548

&lt;211&gt; 415

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 548

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 <212> DNA  
 <213> Pinus radiata

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aagcgcctgt acattgtgct cataagcttg catggtttag taagagggtga aaacatggaa	240
ttgggtagag attctgatac aggcgggcag gttaaatacg ttgtagaact tgcaagagc	299

<210> 550  
 <211> 304  
 <212> DNA  
 <213> Pinus radiata

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 <212> DNA  
 <213> Pinus radiata

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<210> 552  
 <211> 312  
 <212> DNA  
 <213> Eucalyptus grandis

<400> 552	
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<210> 553

<211> 442  
 <212> DNA  
 <213> *Eucalyptus grandis*

<400> 553  
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 cattcttaaa ttccgggaat tg 442

<210> 554  
 <211> 421  
 <212> DNA  
 <213> *Eucalyptus grandis*

<400> 554  
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<210> 555  
 <211> 448  
 <212> DNA  
 <213> *Pinus radiata*

<400> 555  
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 tactgccccaa ctgtgtcctc actgcatcca ccagaatatg gnaaactata aacatttcag 360  
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 gcttaggccc tttgcttcaa aaaaaaaa 448

<210> 556  
 <211> 348  
 <212> DNA  
 <213> *Eucalyptus grandis*

<400> 556  
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 gtatgacagc gtccaggagg aagacttgaa aggttccgg cagtggggaa gcaagacccc 180  
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<210> 557  
 <211> 369  
 <212> DNA  
 <213> *Eucalyptus grandis*

<400> 557  
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<210> 558  
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 <212> DNA  
 <213> *Eucalyptus grandis*

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<210> 559  
 <211> 356  
 <212> DNA  
 <213> *Eucalyptus grandis*

<400> 559  
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 <213> *Eucalyptus grandis*

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 <212> DNA  
 <213> *Eucalyptus grandis*

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 <213> *Eucalyptus grandis*

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 ctggttggga gaaagccctg ccgacgtaca cccagagat tccagcagac gctaccagaa 180  
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 <213> Eucalyptus grandis

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 <213> Eucalyptus grandis

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 <212> DNA  
 <213> Eucalyptus grandis

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 <213> Eucalyptus grandis

<400> 569

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&lt;210&gt; 570

&lt;211&gt; 394

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 570

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&lt;210&gt; 571

&lt;211&gt; 349

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 571

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&lt;210&gt; 572

&lt;211&gt; 388

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 572

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&lt;210&gt; 573

&lt;211&gt; 342

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 573

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&lt;210&gt; 574

&lt;211&gt; 526

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 574

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&lt;210&gt; 575

&lt;211&gt; 295

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 575

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&lt;210&gt; 576

&lt;211&gt; 481

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 576

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&lt;210&gt; 577

&lt;211&gt; 407

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 577

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&lt;210&gt; 578

&lt;211&gt; 332

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 578

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&lt;210&gt; 579

&lt;211&gt; 500

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 579

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&lt;210&gt; 580

&lt;211&gt; 465

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 580

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&lt;210&gt; 581

&lt;211&gt; 494

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 581

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&lt;210&gt; 582

&lt;211&gt; 505

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 582

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&lt;210&gt; 583

&lt;211&gt; 399

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 583

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&lt;210&gt; 584

&lt;211&gt; 472

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 584

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&lt;210&gt; 585

&lt;211&gt; 531

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 585

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&lt;210&gt; 586

&lt;211&gt; 385

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 586

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&lt;210&gt; 587

&lt;211&gt; 314

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 587

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&lt;210&gt; 588

&lt;211&gt; 479

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 588

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&lt;210&gt; 589

&lt;211&gt; 362

&lt;212&gt; DNA



<213> *Eucalyptus grandis*

&lt;400&gt; 589

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ct						362

&lt;210&gt; 590

&lt;211&gt; 190

&lt;212&gt; DNA

<213> *Pinus radiata*

&lt;400&gt; 590

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&lt;211&gt; 301

&lt;212&gt; DNA

<213> *Pinus radiata*

&lt;400&gt; 591

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&lt;210&gt; 592

&lt;211&gt; 468

&lt;212&gt; DNA

<213> *Eucalyptus grandis*

&lt;400&gt; 592

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&lt;210&gt; 593

&lt;211&gt; 601

&lt;212&gt; DNA

<213> *Eucalyptus grandis*

&lt;400&gt; 593

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&lt;210&gt; 594

&lt;211&gt; 239

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 594

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&lt;210&gt; 595

&lt;211&gt; 388

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 595

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&lt;210&gt; 596

&lt;211&gt; 454

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 596

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&lt;210&gt; 597

&lt;211&gt; 443

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 597

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&lt;210&gt; 598

&lt;211&gt; 268

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 598

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&lt;210&gt; 599

&lt;211&gt; 437

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 599

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&lt;211&gt; 578

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 600

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&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

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&lt;210&gt; 610

&lt;211&gt; 399

&lt;212&gt; DNA

<213> *Eucalyptus grandis*

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&lt;210&gt; 611

&lt;211&gt; 363

&lt;212&gt; DNA

<213> *Eucalyptus grandis*

&lt;400&gt; 611

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ctg						363

&lt;210&gt; 612

&lt;211&gt; 457

&lt;212&gt; DNA

<213> *Eucalyptus grandis*

&lt;400&gt; 612

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tggaagtcac	ggctgctcat	ggatgtaaga	agcttggtgt	ttcctcatct	gctactgttt	420
acggttgggc	aaaggagggtc	ccatgtacag	aggactt			457

&lt;210&gt; 613

&lt;211&gt; 383

&lt;212&gt; DNA

<213> *Pinus radiata*

&lt;400&gt; 613

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cgttctggtg	actggaggcg	ctggtttcat	cggaagccac	accgccctgc	agctcctcga	120
ggatggctac	gaggtttata	tcacgcacaa	tttagataac	tctgttgaag	aagcagtga	180

cagagtgagg gatttagttg atcagcgctt ccgcctaaat cttcactttt ttctgggaga	240
tctttgcaac aaagagggat gtagagaagg ttttttcatt ggccaaattc gatgctgtga	300
tacattttgc tggattgaag gctggtggga gaaagtgtag caattccatt acgtaattac	360
caaggaaaaa atctagttgg gca	383

&lt;210&gt; 614

&lt;211&gt; 517

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 614

attcggagaa tttcgctatt gcagaaacca aaacccaaat tggcgatggg gtcacaaggg	60
caaccagagg aaggcttcaa caagtgcgtt ctggtgactg gaggcgctgg tttcatcgga	120
agccacaccg ccctgcagct cctcgaggat ggctacgagg tttatatcat cgacaattta	180
gataactctg ttgaagaagc agtccacaca gtgagggatt tagttgatca gcgcttccgc	240
ctaaatcttc acttttttct gggagatctt tgcaacaaag aggatgtaga gaaggttttt	300
tcattggcca aattcgatgc tgtgatacat tttgctggat tgaaggctgt tggagaaagt	360
gtagcaattc cattacgtta ttacaagaac aatctagttg gcactctgaa cctatatgag	420
attatggcca aacatggttg caaaaagatg gttttttcat catcagctac agtttatggg	480
caaccgaagg gggattccct gtggtagaag actttcc	517

&lt;210&gt; 615

&lt;211&gt; 473

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 615

cattgatatt attttctcac aaaaaaaagt ccgggttatg tcaagaagat ctgctgttta	60
ttcctcacca tgcagatccc ctgaatcaac taatctagaa tcgtcatttg ggtattttatt	120
gaggcatggt actctttaga tttttgaagt ttttgggcta atacagagcg atagttagca	180
tggagtgccca aggaaagaac attctggtca ccggaggagc aggttatggt ggcagtcaca	240
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cttcagaaga agctattaca agagttgcta agctcgctgg cgaatatggg ggcaatctca	360
ccttccataa gattgatctt ctgggtaaaa gaagctatgg agaaattggt cttatcaaca	420
gaatttgatg ctgtcattca ttttctggtg gttaaagctg tcggagagag tgt	473

&lt;210&gt; 616

&lt;211&gt; 323

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 616

actttctgct aatgaatcca tatgggacgc accaagctct tccttgaaga gattagcccg	60
tgatatttat aaggcagatc cagattggag aataattttg ctgaggtact tcaaccaggt	120
tggagctcac ccaagtggcc agattggtga agatccaaag ggaattccaa ataacctcat	180
gcctttcatc caacaagtgg ctgtgggaag gcaaccagtg ctgaacgtat atggtaatga	240
ttaccaaaaca aaggatggca cagcggttcg agattacatt catgtggtag acttggctga	300
tggtcacatt tctgcgcttc aga	323

&lt;210&gt; 617

&lt;211&gt; 497

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 617

gagaatttctg ctattgcaga aacccaaacc aaaattggcg atgggggtcac aagggaacc	60
agaggaaggc ttcaacaagt gcgttctggt gactggaggc gctggtttca tcggaagcca	120

caccgccctg	cagctcctcg	aggatggcta	cgaggtttat	atcatcgaca	atthagataa	180
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aattccatta	cgttattaca	agaacaatct	agttggcact	ctgaacctat	atgagattat	420
ggccaacatg	gttgcaaaaa	gatggttttt	tcatcatcag	ctacagttat	ggccaaccca	480
aggtgattcc	ttgtgta					497

&lt;210&gt; 618

&lt;211&gt; 384

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 618

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atggggggcaa	tctcaccttc	cataagattg	atcttctgga	taaagaagct	atggagaaat	240
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agagtgtagc	aaagccactg	ctttactaca	aaaacaacat	agttggcacc	ttaaacttat	360
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&lt;210&gt; 619

&lt;211&gt; 354

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 619

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atggggggcaa	tctcaccttc	cataagattg	atcttctgga	taaagaagct	atggagaaat	240
tggtcttctc	aacagaattt	gatgctgtca	ttcattttgc	tgggttgaaa	gctgtcggag	300
agagtgtagc	aaagccactg	ctttactaca	aaaacaacat	agttggcacc	ttaa	354

&lt;210&gt; 620

&lt;211&gt; 425

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 620

gaaggcttca	acaagtgcgt	tctggtgact	ggaggcgctg	gtttcatcgg	aagccacacc	60
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ccattacggt	attacaagaa	caatctagtt	ggcactctga	acctatatga	gattatggcc	360
aaacatgggt	gcaaaaagat	ggttttttca	tcatcagcta	cagtttatgg	ccaacccaag	420
gtgat						425

&lt;210&gt; 621

&lt;211&gt; 623

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 621

ggaatcagag	aaatggctgc	acacgcagtg	gctctggttc	tgggtcttgt	tctgatggct	60
------------	------------	------------	------------	------------	------------	----



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ggaccttggg	gtgaaattcc	ccttcaacca	gccgatgaaa	ttgtactcca	gcctgtggaa	600
tgcggtatgct	gggcactcgg	gga				623

&lt;210&gt; 622

&lt;211&gt; 426

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 622

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atcaagctcg	tcgagggcga	ctcggctggg	accgtcactg	ctttctacat	gtcgtcggat	360
ggaccgaacc	acaacgaatt	cgacttcgag	ttcctgggca	acacgacagg	ggagccctac	420
ctggtc						426

&lt;210&gt; 623

&lt;211&gt; 412

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 623

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gggtcttgct	ttgttgggtg	gtctggctgc	gggtgcgagg	tttgaggagc	tctaccagcc	180
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gatcaagctc	gtcagggcgc	actcggctgg	gaccgtcact	gctttctaca	tgtcgtcgga	360
tggaccgaac	cacaacgaat	tcgacttcga	gtcctgggca	aacgacaggg	ga	412

&lt;210&gt; 624

&lt;211&gt; 373

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 624

gcgccaagtc	gtgtttcttg	tggacgagac	accgatccgc	gtgcacacca	acatggcgca	60
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cacggtgtcc	gagctgagcc	tccaccagaa	ccaccagctc	aagtgggtcc	aggcccanca	360
catggtctac	gac					373

&lt;210&gt; 625

&lt;211&gt; 351

&lt;212&gt; DNA

## &lt;213&gt; Eucalyptus grandis

## &lt;400&gt; 625

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tcctactctg	ttctctggaa	catgtaccag	attgtattct	tcgtggatga	cgtgccgac	120
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gactggtcca	aggcgccgtt	cgtggcctct	taccgggggt	tccacattga	cgggtgcgaa	300
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## &lt;210&gt; 626

## &lt;211&gt; 270

## &lt;212&gt; DNA

## &lt;213&gt; Eucalyptus grandis

## &lt;400&gt; 626

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gatgactggg	ccactcgggg	agggcttgag				270

## &lt;210&gt; 627

## &lt;211&gt; 267

## &lt;212&gt; DNA

## &lt;213&gt; Eucalyptus grandis

## &lt;400&gt; 627

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cttgggggtg	aattccccct	caaccagccc	atgaaattgt	actccagcct	gtggaatgcg	240
gatgactggg	ccactcggga	ggcttga				267

## &lt;210&gt; 628

## &lt;211&gt; 468

## &lt;212&gt; DNA

## &lt;213&gt; Eucalyptus grandis

## &lt;400&gt; 628

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gtcgtcggat	ggaccgaacc	acaacgaatt	cgacttcgag	ttcctgggca	acacgacagg	420
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## &lt;210&gt; 629

## &lt;211&gt; 559

## &lt;212&gt; DNA

## &lt;213&gt; Eucalyptus grandis

## &lt;400&gt; 629

gagaaacaga	gcgacttcag	agctcgtgaa	gaaaagcttt	tgctctcgct	cttcttcttc	60
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ggctcggcct	ctgggtcgac	ccccaccact	gacttccact	cctactccgt	cctctggaac	540
caaccgccaa	agtcgtgtt					559

&lt;210&gt; 630

&lt;211&gt; 416

&lt;212&gt; DNA

<213> *Eucalyptus grandis*

&lt;400&gt; 630

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ctgtcccggg	cttttccaaa	gtgtctgtgt	cgttcggcct	gttcgtgggt	cttgcttgt	120
tggtgggtct	ggtcgcgggt	gcaaggttta	aggagctcta	ccagccgggc	tgggctatgg	180
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acgaattcga	cttcgagttc	ctggggcaac	acgacagggg	aacctacct	ggtaca	416

&lt;210&gt; 631

&lt;211&gt; 250

&lt;212&gt; DNA

<213> *Eucalyptus grandis*

&lt;400&gt; 631

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aggaccagcc	catgggcgtc	tacagctcga	tatggaacgc	cgacgactgg	gccaccagg	120
gcggccgcat	caagaccgac	tggaccacag	cccccttcac	cacgtcctac	cgtaacttcg	180
agatcgacgc	gtgcgagtgc	ccggcgacaa	tggcgccggc	agacaccgcc	aagcggtgca	240
gcagcgccgg						250

&lt;210&gt; 632

&lt;211&gt; 475

&lt;212&gt; DNA

<213> *Eucalyptus grandis*

&lt;400&gt; 632

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ttcttcttct	tcttcttcaa	tggtgtccc	ggtcttttcc	aaagtgtctg	tgctgttcgg	120
cctgttcgtg	ggtcttgcgt	tggtgggtggg	tctgggtcgcg	ggtgcgaggt	ttgaggagct	180
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gtcgtcggat	ggaccgaacc	acaacgaatt	cgacttcgag	ttcctgggaa	cacgacaggg	420
gatccctacc	tggtacagac	caacgtgtac	gtgaacgggg	tgggaaccgg	gacaa	475

&lt;210&gt; 633

&lt;211&gt; 416

&lt;212&gt; DNA

<213> *Eucalyptus grandis*

&lt;400&gt; 633

gagcgacttc	agagctcgtg	aagaaaagct	tttgcctcgc	ctcttcttct	tcttcttctt	60
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tgggtcttgc	gttggttggtg	ggtctgggtc	cgggtgctgag	gtttgaggag	ctctaccagc	180
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atggaccgaa	ccacaacgaa	ttcgacttcg	agttcctggtg	caacacgaca	ggggag	416

&lt;210&gt; 634

&lt;211&gt; 232

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 634

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tcctactctg	ttctctggaa	catgtaccag	attgtattct	tcgtggatga	cgtgccgatc	120
cgggtgttca	agaacagcaa	ggaccttggg	gtgaaattcc	ccttcaacca	gccgatgaaa	180
ttgtactcca	gcctgtggaa	tgcggatgac	ttgggcaatc	cggggagggt	tt	232

&lt;210&gt; 635

&lt;211&gt; 287

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 635

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tgaggccaag	tgctgcgcta	ctcagggcca	gaggtgggtg	gaccagaagg	agttccagga	180
cctcgatgcc	ttccagtacc	ggaggctccg	gtgggtgcgc	tcgagataca	ccatctacaa	240
ctactgcgct	gatcggnaga	ggtaccccg	gatntccccg	gagtgc		287

&lt;210&gt; 636

&lt;211&gt; 240

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 636

ggacgagaca	ccgatccgcg	tgcacaccaa	cttgagacac	cggggcatcc	cgtacccgaa	60
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&lt;210&gt; 637

&lt;211&gt; 360

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 637

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&lt;210&gt; 638

&lt;211&gt; 401

&lt;212&gt; DNA

## &lt;213&gt; Eucalyptus grandis

&lt;400&gt; 638

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&lt;210&gt; 639

&lt;211&gt; 461

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 639

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&lt;210&gt; 640

&lt;211&gt; 458

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 640

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&lt;210&gt; 641

&lt;211&gt; 283

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 641

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&lt;210&gt; 642

&lt;211&gt; 385

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 642

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&lt;210&gt; 643

&lt;211&gt; 378

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 643

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&lt;210&gt; 644

&lt;211&gt; 430

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 644

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&lt;210&gt; 645

&lt;211&gt; 471

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 645

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&lt;210&gt; 646

&lt;211&gt; 480

&lt;212&gt; DNA

## &lt;213&gt; Eucalyptus grandis

&lt;400&gt; 646

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&lt;210&gt; 647

&lt;211&gt; 284

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 647

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&lt;210&gt; 648

&lt;211&gt; 459

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 648

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&lt;210&gt; 649

&lt;211&gt; 402

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 649

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&lt;210&gt; 650

&lt;211&gt; 469

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

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&lt;210&gt; 651

&lt;211&gt; 473

&lt;212&gt; DNA

<213> *Eucalyptus grandis*

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&lt;210&gt; 652

&lt;211&gt; 454

&lt;212&gt; DNA

<213> *Eucalyptus grandis*

&lt;400&gt; 652

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&lt;210&gt; 653

&lt;211&gt; 435

&lt;212&gt; DNA

<213> *Eucalyptus grandis*

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 <213> Eucalyptus grandis

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&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 658

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&lt;210&gt; 659

&lt;211&gt; 482

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 659

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&lt;210&gt; 660

&lt;211&gt; 415

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 660

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&lt;210&gt; 661

&lt;211&gt; 542

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 661

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 tatg 424

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<400> 668  
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 <213> Eucalyptus grandis

<400> 669  
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<212> DNA  
<213> Eucalyptus grandis

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<212> DNA  
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<212> DNA  
<213> Eucalyptus grandis

<400> 672  
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<213> Eucalyptus grandis

<400> 673  
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<210> 674

&lt;211&gt; 387

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 674

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&lt;210&gt; 675

&lt;211&gt; 324

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 675

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gcgttccata	tcgagctgta	gacg				324

&lt;210&gt; 676

&lt;211&gt; 330

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 676

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&lt;210&gt; 677

&lt;211&gt; 438

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 677

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&lt;210&gt; 678

&lt;211&gt; 362

&lt;212&gt; DNA

## &lt;213&gt; Eucalyptus grandis

## &lt;400&gt; 678

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ac						362

## &lt;210&gt; 679

## &lt;211&gt; 424

## &lt;212&gt; DNA

## &lt;213&gt; Eucalyptus grandis

## &lt;400&gt; 679

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tgac						424

## &lt;210&gt; 680

## &lt;211&gt; 414

## &lt;212&gt; DNA

## &lt;213&gt; Eucalyptus grandis

## &lt;400&gt; 680

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## &lt;210&gt; 681

## &lt;211&gt; 239

## &lt;212&gt; DNA

## &lt;213&gt; Eucalyptus grandis

## &lt;400&gt; 681

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## &lt;210&gt; 682

## &lt;211&gt; 319

## &lt;212&gt; DNA

## &lt;213&gt; Eucalyptus grandis

## &lt;400&gt; 682

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&lt;210&gt; 683

&lt;211&gt; 424

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 683

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ggcc						424

&lt;210&gt; 684

&lt;211&gt; 309

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 684

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acgtgaacg						309

&lt;210&gt; 685

&lt;211&gt; 238

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 685

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&lt;210&gt; 686

&lt;211&gt; 515

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 686

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&lt;210&gt; 687

&lt;211&gt; 445

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 687

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&lt;210&gt; 688

&lt;211&gt; 422

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 688

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ca						422

&lt;210&gt; 689

&lt;211&gt; 279

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 689

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&lt;210&gt; 690

&lt;211&gt; 452

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 690

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&lt;210&gt; 691

&lt;211&gt; 346

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 691

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&lt;210&gt; 692

&lt;211&gt; 470

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 692

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&lt;210&gt; 693

&lt;211&gt; 374

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 693

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&lt;210&gt; 694

&lt;211&gt; 409

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 694

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&lt;210&gt; 695

&lt;211&gt; 224

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 695

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&lt;210&gt; 696

&lt;211&gt; 442

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 696

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&lt;210&gt; 697

&lt;211&gt; 408

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 697

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&lt;210&gt; 698

&lt;211&gt; 469

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 698

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<210> 699

<211> 347

<212> DNA

<213> Eucalyptus grandis

<400> 699

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<210> 700

<211> 452

<212> DNA

<213> Eucalyptus grandis

<400> 700

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<210> 701

<211> 323

<212> DNA

<213> Eucalyptus grandis

<400> 701

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<210> 702

<211> 441

<212> DNA

<213> Eucalyptus grandis

<400> 702

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<400> 705  
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<400> 706  
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&lt;211&gt; 397

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 716

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&lt;210&gt; 717

&lt;211&gt; 365

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 717

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&lt;210&gt; 718

&lt;211&gt; 301

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 718

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a						301

&lt;210&gt; 719

&lt;211&gt; 383

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 719

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&lt;210&gt; 729

&lt;211&gt; 538

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 729

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&lt;210&gt; 730

&lt;211&gt; 412

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 730

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&lt;210&gt; 731

&lt;211&gt; 350

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 731

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&lt;210&gt; 732

&lt;211&gt; 354

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 732

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&lt;210&gt; 733

&lt;211&gt; 480

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 733

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&lt;210&gt; 734

&lt;211&gt; 343

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 734

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&lt;210&gt; 735

&lt;211&gt; 359

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 735

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&lt;210&gt; 736

&lt;211&gt; 360

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 736

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&lt;210&gt; 737

&lt;211&gt; 437

&lt;212&gt; DNA

<213> *Eucalyptus grandis*

&lt;400&gt; 737

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&lt;210&gt; 738

&lt;211&gt; 341

&lt;212&gt; DNA

<213> *Eucalyptus grandis*

&lt;400&gt; 738

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&lt;210&gt; 739

&lt;211&gt; 497

&lt;212&gt; DNA

<213> *Eucalyptus grandis*

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&lt;210&gt; 740

&lt;211&gt; 497

&lt;212&gt; DNA

<213> *Eucalyptus grandis*

&lt;400&gt; 740

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&lt;210&gt; 741

&lt;211&gt; 395

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 741

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&lt;210&gt; 742

&lt;211&gt; 396

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 742

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&lt;211&gt; 347

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 743

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&lt;211&gt; 446

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 744

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&lt;211&gt; 439

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 745

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&lt;210&gt; 746

&lt;211&gt; 322

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 746

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&lt;210&gt; 747

&lt;211&gt; 433

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 747

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&lt;211&gt; 525

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 748

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&lt;210&gt; 749

&lt;211&gt; 385

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 749

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&lt;211&gt; 519

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 750

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&lt;210&gt; 751

&lt;211&gt; 342

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 751

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&lt;210&gt; 752

&lt;211&gt; 416

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis



&lt;400&gt; 752

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&lt;210&gt; 753

&lt;211&gt; 408

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 753

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&lt;213&gt; Eucalyptus grandis

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&lt;211&gt; 403

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 755

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&lt;210&gt; 756

&lt;211&gt; 414

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 756

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&lt;211&gt; 441

&lt;212&gt; DNA

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&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

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&lt;211&gt; 340

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

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&lt;210&gt; 769

&lt;211&gt; 368

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 769

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&lt;210&gt; 770

&lt;211&gt; 342

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 770

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&lt;210&gt; 771

&lt;211&gt; 580

&lt;212&gt; DNA

&lt;213&gt; Eucalyptus grandis

&lt;400&gt; 771

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&lt;211&gt; 407

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 772

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&lt;210&gt; 773

&lt;211&gt; 403

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 773

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&lt;210&gt; 774

&lt;211&gt; 400

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 774

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&lt;211&gt; 384

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 775

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&lt;210&gt; 776

&lt;211&gt; 345

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 776

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&lt;210&gt; 777

&lt;211&gt; 449

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 777

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&lt;210&gt; 778

&lt;211&gt; 354

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 778

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&lt;210&gt; 779

&lt;211&gt; 392

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 779

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&lt;210&gt; 780

&lt;211&gt; 293

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 780

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&lt;211&gt; 442

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 790

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&lt;210&gt; 791

&lt;211&gt; 424

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 791

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&lt;210&gt; 792

&lt;211&gt; 219

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 792

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&lt;210&gt; 793

&lt;211&gt; 405

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 793

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<212> DNA  
<213> Pinus radiata

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 <212> DNA  
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<400> 801  
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 <212> DNA  
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<400> 802  
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&lt;210&gt; 803

&lt;211&gt; 429

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 803

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&lt;210&gt; 804

&lt;211&gt; 432

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 804

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&lt;210&gt; 805

&lt;211&gt; 438

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 805

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&lt;210&gt; 806

&lt;211&gt; 393

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

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 <212> DNA  
 <213> Pinus radiata

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 catatttgtt tggaaaagca ggggcacaga ttaagctcgt tccagggtgac tctgcaggca 180  
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<210> 808  
 <211> 440  
 <212> DNA  
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 gggctactca agggggcctc gtcaagaccg attggagcca cgcacctttc atttccacat 180  
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 <213> Pinus radiata

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 agctggggct cagatcacat caaacaattc catggcggtc gaaagactga gctgctcctc 180  
 aacaaacagt atggtgcggg gtttgantcg aaggggacat atttatttgg gcatttcagt 240  
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<210> 810  
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ggt						423

&lt;210&gt; 811

&lt;211&gt; 483

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 811

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&lt;210&gt; 812

&lt;211&gt; 323

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 812

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&lt;210&gt; 813

&lt;211&gt; 430

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 813

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caagagcaag						430

&lt;210&gt; 814

&lt;211&gt; 331

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 814

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&lt;210&gt; 815

&lt;211&gt; 257

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 815

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&lt;210&gt; 816

&lt;211&gt; 216

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 816

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gagggggtct	ggagaaaacc	gactgggcca	aggcgc			216

&lt;210&gt; 817

&lt;211&gt; 393

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 817

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&lt;210&gt; 818

&lt;211&gt; 457

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 818

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<210> 820  
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 <212> DNA  
 <213> Pinus radiata

<400> 820  
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<400> 822  
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<210> 823  
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<213> Pinus radiata

<400> 823

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<210> 824

<211> 328

<212> DNA

<213> Pinus radiata

<400> 824

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tggtagtcca	tgtacttcgt	gggtgggatca	gccggcgtag	gcctcactta	ccgcaaagca	300
gatcatgcaa	ctcaagtggg	tacacgag				328

<210> 825

<211> 352

<212> DNA

<213> Pinus radiata

<400> 825

gcatttacct	ctgggtcgac	cccaccgcag	atttccattc	ctattctttt	ctgtggaacc	60
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ggctgggagt	cccatatcct	aagaaacagc	ccatgagggt	atcctcttca	atctggaatg	180
cagataactg	ggctactcaa	gggtggcggc	tgaagataaa	ctggagccat	tctcctttta	240
tctccactta	caaaaagggtc	gacatcgatg	caaaccaata	cggattaaat	ggagaatcaa	300
gagggtttat	tgagaatgga	agtaagtggg	gggacaggcc	ctctcattct	tc	352

<210> 826

<211> 215

<212> DNA

<213> Pinus radiata

<400> 826

gccagctggg	gctcagatca	catcaaacia	ttccatggcg	gtcgaaagac	tgagctgctc	60
ctcaacaaac	agtatgggtg	ggggtttgag	tcgaagggga	catatttatt	tgggcatttc	120
agtatgcaga	taaagctggg	tgccggtgat	tccgctggca	ctgtcaccgc	ctttnatctt	180
tcttctcaga	ctgcagagca	cgatgagatg	acttt			215

<210> 827

<211> 463

<212> DNA

<213> Pinus radiata

<400> 827

gttgacatgg	cttgtttaag	attgcagagc	tgctgctttt	tcgttctggg	ttttttgctt	60
ctgggtatct	aattgtgcag	agttcaatga	tatcttcgag	cccagctggg	cgattgatca	120
tgttatgaac	gagggagagc	tggtgaagct	gaagcttggc	cattttttct	ggcgctggct	180
tttcttccaa	ggccacatac	ttgtttggaa	aagttagggc	gcagattaaa	ctcgttcccc	240

gcgactctgc	gggcacagtg	actgcatttt	atatgtcttc	tgaggggaca	ttgcatgacg	300
aattcgattt	cgaattcttg	ggaaatgctt	cgggtgagcc	ttacattgtg	cagactaata	360
tctactccaa	cggcactggc	gacagggaac	aacgcattta	cctctgggtc	gaccccaccg	420
cagatttcca	ttcctattct	tttctgtgga	accacaagca	agt		463

&lt;210&gt; 828

&lt;211&gt; 342

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 828

caacttttct	ggcgctggct	tttcttccaa	ggcaacatac	ttgtttggaa	aagtaggggc	60
gcagattaaa	ctcgttcccg	gcgactctgc	gggcacagtg	actgcatttt	atatgtcttc	120
tgaggggaca	ttgcatgacg	aattcgattt	cgaattcttg	ggaaatgctt	cgggtgagcc	180
ttacattgtg	cagactaata	tctactccaa	tggcactggc	aacagggaac	aacgcattta	240
cctctgggtc	gaccccaccg	cagatttcca	ttcctattct	tttctgtgga	accacaagca	300
agttgtattc	tttgtagaca	gtgttccgat	tagggtattc	cc		342

&lt;210&gt; 829

&lt;211&gt; 447

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 829

ccggtctggt	gttgtgggta	ataggatggc	cgggcaaagg	aattggttca	agagaatcga	60
gtttattggt	atattcgtgg	tttgcttaaa	ctctgtttct	gcacgcccgg	catcatttgc	120
agaggatttt	aaagtgcgt	gggcagatga	ccatgtcaaa	acaaggtcag	ataacaactc	180
catcgatctc	atcctggatc	agaattcagg	ggcaggattt	gcctccaaga	atcagtacat	240
gtttggactt	gtaagcatga	acatcaaact	tgtggcgggt	gattctgcag	ggacagtcac	300
tgctttttat	atgagctcgg	acaaggagga	agtgcgagat	gaattggatt	tcgagtttct	360
ggggaacaga	tcaggccagc	cttatacagt	ccaaacaaat	gtgtttgctc	tcgggaaggg	420
tggccgcgag	cagagagtga	atctctg				447

&lt;210&gt; 830

&lt;211&gt; 471

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 830

tgtttaagaa	tgacagattg	ctgcttcttc	attctgggtt	tttgcttctg	ggtatctaata	60
tgtgcagagt	tcaatgatat	cttcgagccc	agctgggcga	ttgatcatgt	tatgaacgag	120
ggagagctgt	tgaagctgaa	gctcgacaac	ttttctggcg	ctggcttttc	ttccaaggca	180
acatacttgt	ttggaaaagt	aggggcgcag	attaaactcg	ttcccggcga	ctctgcgggc	240
acagtgactg	cattttatat	gtcttctgag	gggacattgc	atgacgaatt	cgatttcgaa	300
ttcttgggaa	atgcttcggg	tgagccttac	attgtgcaga	ctaatatcta	ctccaatggc	360
actggcaaca	gggaacaacg	catttacctc	tggttcgacc	ccaccgcaga	tttccattcc	420
tattcttttc	tgtggaacca	caagcaagtt	gtattctttg	tagacagtgt	t	471

&lt;210&gt; 831

&lt;211&gt; 391

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 831

taggatggcc	gggcaaaagga	attggttcaa	gagaatcgag	tttattgtta	tattcgtggt	60
ttgtctaaac	tctgtttctg	cacgcccgcc	atcattttga	gaggatttta	aagtgcagtg	120
ggcagatgac	catgtcaaaa	caaggtcaga	taacaactcc	atcgatctca	tcctggatca	180

gaattcaggg	gcaggatttg	cctccaagaa	tcagtacatg	tttggacttg	taagcatgaa	240
catcaaaactt	gtggcgggtg	attctgcagg	gacagtcact	gctttttata	tgagctcggg	300
caaggaggaa	gtgcgagatg	aattggattt	cgagtttctg	gggaacagat	caggccagcc	360
ttatacagtc	caaacaaatg	tgtttgctct	c			391

&lt;210&gt; 832

&lt;211&gt; 304

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 832

catccctgc	tttcgagga	acacatggcg	tgtagcctta	gtgtgcctgt	ggactctcaa	60
tggaacttg	tccacgaaaa	acacaatctg	atgagcgttc	cagaggatag	agtaagtgtg	120
aaactctgtc	gttggatcga	accagaggat	atgcctctgt	tcccttcac	ccacaccatt	180
tgaatagatg	ttcgtctgaa	gaccataagg	ctgtccactt	ctgttcctta	gaaattcaaa	240
atctagctcg	tccctattca	tgtctgtgtc	agaagacata	taataagcag	tgacaactcc	300
tgca						304

&lt;210&gt; 833

&lt;211&gt; 234

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 833

acccatttaa	ccagccgatg	aagatctatt	cgagcctgtg	gaatgctgat	gactgggcca	60
cccagggggg	tctggagaaa	accgactggg	ccaaggcgcc	cttcacgccc	tccacagggg	120
aattccacgt	cgatgcctgt	gaggcttctg	ctccgcaatc	ggtgtgcgct	acgcaggggc	180
ggnggtggtg	ggatcaggag	gagttcagag	acctggatgg	gcggcaatgg	cggt	234

&lt;210&gt; 834

&lt;211&gt; 375

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 834

ggagcattcg	cctttgtaat	ggagtaaggg	aagaatgaga	gggcctgtcc	caccacttac	60
ttccattttt	aataacccct	cttggttctc	catttaatcc	gtatttggtt	gcatcaatgt	120
tgaagccttt	gtaagtggag	ataaaaaggag	aatggctcca	gtttatcttc	agccgcccac	180
cttgagtagc	ccagttatct	gcattccaga	ttgaagagga	taccctcatg	ggctgtttct	240
taggatatgg	gactcccagc	ctctcgttgt	tggggaatac	cctaatacga	acactgtcta	300
caaagaatac	aacttgcttg	tggttcaca	gaaaagaata	ggaatggaaa	tctgcggtgg	360
ggtcgaacca	gaggt					375

&lt;210&gt; 835

&lt;211&gt; 352

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 835

gcagctttct	cttgacaaat	ggacagggtac	tggcttccaa	tccaagggtg	gctacttggt	60
tggacatttc	agtatgcaga	ttaaagatgg	tcctgggtgac	tctgcaggcg	ttgtgactgc	120
cttttattta	tcctctcaga	actctgaaca	tgatgaaata	gactttgagt	tcttgggcaa	180
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ttggaatatg	catcagattg	tattctttgt	ggatgatgtc	cccatcagag	tt	352

&lt;210&gt; 836

<211> 368  
 <212> DNA  
 <213> Pinus radiata

<400> 836  
 ctataagtc ggattcttca gtgctgctat taagctacag gcaggttata cagctggagt 60  
 cattgcagca ctctatctct ccaataacca ggagtaccca ggtcaccatg acgaaataga 120  
 cattgagttc ctggggacaa caccaggaaa accctacacc ttacagacca atgtttacat 180  
 aaatggaaca ggggatgggc aggttctcac aggcagggag ttgaagtttc atctctggtt 240  
 tgacccaact gaagacttcc ataactacag ccttctctgg actccaagtt atatcatctt 300  
 ctatgtagat gatattgcta tccgaaagta cccaagaaga atttcatcta cttatccatt 360  
 gaggccac 368

<210> 837  
 <211> 402  
 <212> DNA  
 <213> Pinus radiata

<400> 837  
 ggagcattcg cctttgtaat ggagtaaggg aagaatgaga gggcctgtcc caccacttac 60  
 ttccattttt aataaccctt cttgggttctc catttaatcc gtatttgttt gcatcaatgt 120  
 tgaagccttt gtaagtggag ataaaaggag aatggctcca gtttatcttc agccgcccac 180  
 cttgagtagc ccagttatct gcattccaga ttgaagagga taccctcatg ggctgtttct 240  
 taggatattg gactcccagc ctctcgttgt tggggaatac cctaatacgga aactgtctta 300  
 caaagaatac aacttgcttg tggttccaca gaaaagaata ggaatggaaa tctgcggtgg 360  
 ggtcgaacca gaggtaaatg cgttggtccc tgtcgccagt gc 402

<210> 838  
 <211> 389  
 <212> DNA  
 <213> Pinus radiata

<400> 838  
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 aatgctgac aatatgggtc ccaatgtact gttatttctc ttggtagcgg caatggctgc 120  
 tactgctacc tcacctcga agcctgtgga tgtgccattc caaaaaaact atgtaccac 180  
 ctgggcttct gatcatatca agtacattaa tggggggaac gaagcgcagc tttctcttga 240  
 caaatggaca ggtactggct tccaatccaa gggtagctac ttgtttggac atttcagtat 300  
 gcagataaag atgggttcctg gtgactctgc aggcgttgtg actgcctttt atttatcctc 360  
 tcagaactct gaacatgatg aaatagact 389

<210> 839  
 <211> 451  
 <212> DNA  
 <213> Pinus radiata

<400> 839  
 ggatttagga gtgaggtatc cattcaacca gcccatgaaa atctattcaa gcttggtgaa 60  
 tgctgatgac tgggctacaa ggggtgggtt ggagaagaca gactggagca aggcaccctt 120  
 tgttgcatca tacaaggat tccacgtgga tgggtgtgag gcgtctatgc ctactctgc 180  
 ttgtccaact ttaggccgtc gatggtggga tcagaaagcc ttcatgacc ttgatggaca 240  
 gcaatggagg aaactgaagt ggggttcgtga taggtacacc atatacaact actgcactga 300  
 cagagtggag tatectaaaa tgtctccaga gtgtaccana gaccgtgaca tctaatagca 360  
 cagcctcctt gggatagata gctatatttt tattctattc ttcttcgaca tatggctgtt 420  
 ctaattatgt tataactgcc atttcgtagt a 451

<210> 840

<211> 459  
 <212> DNA  
 <213> Pinus radiata

<400> 840  
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 ctgctattaa gctacaggca ggttatacag ctggagtcac tgcagcactc tatctctcca 120  
 ataaccaggga gtacccagggt caccatgacg aaatagacat tgagtccctg gggacaacac 180  
 caggaaaacc ctacacctta cagaccaatg tttacataaa tggaaacaggg gatgggcagg 240  
 ttctcacagg cagggagttg aagtttcatc tctggtttga cccaactgaa gacttccata 300  
 actacagcct tctctggact ccaagttata tcatcttcta tgtagatgat attgctatcc 360  
 gaaagtaccc aagaagaatt tcatctactt atccattgag gccactttgg gtatatggat 420  
 caatatggga tgcttcctct tgggctactg aaaatggca 459

<210> 841  
 <211> 476  
 <212> DNA  
 <213> Pinus radiata

<400> 841  
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 gattgatacc ctctaataatt ttacatgtgt gtctgagaaa gaatgaaaga attaatgtaa 120  
 tagaaatgaa tcagaataact tggaggaaa gaattcagag ttcattctct gaattcaaag 180  
 gattgaggtt ttatttgcac tggggaggcg gtccctgggg gaatctcttg gtgtctgcac 240  
 aataatcgta gatcatgtag ttctttcgta cccactgtag cctctgcctc tgtcttgcct 300  
 tcgattccaa agccacctcc tctgcccggc cgtcaaaaaga agaagtagcg gcagcattga 360  
 aattgcggaa ggatgcaaca aaggagcgtt tagtccagtc ggtcttcaca agccccccc 420  
 tggttgcccc atcatctgca ttccaaaggc tggagtagac cctcattgct ttgggt 476

<210> 842  
 <211> 293  
 <212> DNA  
 <213> Pinus radiata

<400> 842  
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 ctctgtttct tcgtcagagt ccgcgcctcg ggcttatccc accagtgcct tgcacttgca 120  
 ttattactgt gagcacagga cgattgggtc gtgttttgat cctcccatct gcagccgagg 180  
 ccatggaatt tcttgtacga agatacgaac ggagcagccg cccagttggt cttctcaagg 240  
 ccgcctctcg ttgcccagtt gtctccattc caaatgctgg agaactgta cat 293

<210> 843  
 <211> 460  
 <212> DNA  
 <213> Pinus radiata

<400> 843  
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 aagggttttg atgataattt tcagatattg tgggctcagg atcacttcag gacctctgaa 120  
 aatggtcaag tatggcacct ggttcttgac cagaactcag gttctgggtt caaatcgaag 180  
 aacaagtata gattcggatg gtccagcatg aagctcaagc tcgtaccagg agactctgca 240  
 ggagttgtca ctgcttatta tatgtcttct gacacagaca tgaataggga cgagctagat 300  
 tttgaatttc tagggaacag aagtggacag ccttatggtc ttcagacgaa catctattca 360  
 aatggtgtgg gtggaaggga acagaggcat atcctctggt tcgatccanc gacagagttt 420  
 tacacttact ctatcctctg gaacgctcat cagattgtgt 460

<210> 844

<211> 491  
 <212> DNA  
 <213> Pinus radiata

<400> 844  
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 tgaaatagac tttgagttct tgggcaatag gtctggacaa ccttacattc tccaaactaa 120  
 tgttttcagt ggaggaaagg gggacagaga gcaacgcgtat tatctctggt ttgacccac 180  
 aaaagactat cattcctaca ctgtcctttg gaatatgcat cagattgtat tctttgtgga 240  
 tgatgtcccc atcagagttt tcaagaacag caaggattta ggagtggagt atccattcaa 300  
 ccagcccatg aaaatctatt caagcttggt gaatgctgat gactgggcta caaggggtgg 360  
 gttggagaag acagactgga gcaaggcacc ctttgttgca tcatacaagg gattccacgt 420  
 ggatgggtgt gaggcgtcta tgcctcactc tgcttgctca actttaggcc cgtcgtggt 480  
 gggatcaaga a 491

<210> 845  
 <211> 413  
 <212> DNA  
 <213> Pinus radiata

<400> 845  
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 taggcaatgc tgattaatat ggtgccaat gtactgttat ttctcttggt agcggcaatg 120  
 gctgctactg ctaccccacc tccgaagcct gtggatgtgc cattccaaaa aaactatgta 180  
 cccacctggg cttctgatca tatcaagtac attaatgggg ggaacgaagc gcagctttct 240  
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 agtatgcaga taaagatggg tcctgggtgac tctgcaggcg ttgtgactgc cttttattta 360  
 tcctctcaga actctgaaca tgatgaaata gactttgagt tcttgggcaa tag 413

<210> 846  
 <211> 513  
 <212> DNA  
 <213> Pinus radiata

<400> 846  
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 ggggtctactc cagccttttg aatgcagatg attgggcaac caggggtggg cttgtgaaga 120  
 ccgactggac taaagctccc tttgttgcat ccctccgcaa tttcaatgct gccgctactt 180  
 cttcttttga tgccgtcgca gaggaggtgg ctttgggaatc gaaccaagaa cagaggcaga 240  
 ggctccagtg ggtacgaaag aactacatga tctacgatta ttgtgcagac accaagagat 300  
 tccccaggg accgcctccc gattgcaaat aaaacctcaa tcctttgaat tcagagaatg 360  
 aactctgaat tctaccttcc aagtattctg attcatttct attacattaa ttctcattct 420  
 ttctcagaca cacatgtaaa atattagagg gtatcaatca tttcattttg gtactgctat 480  
 acaaaagcga gattcttttg atgaaaaaaa aaa 513

<210> 847  
 <211> 362  
 <212> DNA  
 <213> Pinus radiata

<400> 847  
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 cagggagttg aagtttcatc tctggtttga cccaactgaa gacttccata actacagcct 120  
 tctctggact ccaagttata tcatcttcta tgtagatgat attgctatcc gaaagtaccc 180  
 aagaagaatt tcatctactt atccattgag gccacttttg gtatatggat caatatggga 240  
 tgcttctctc tgggctactg aaaatggcaa atacagagca gattacagat atcagccatt 300  
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ag

362

<210> 848  
 <211> 417  
 <212> DNA  
 <213> Pinus radiata

<400> 848

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gcagctcttg	cttctgcaaa	tttctacaat	gatgtcgaca	ttacatgggg	taatgatcgt	180
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ggatttcaat	ccaagcaaga	gtatctattt	gccaagattg	atatccaaat	caagttggta	300
cctggcaact	ctgcaggcac	agtcactacc	ttttatctat	catctcaagg	tcccaaacac	360
gacgaaatag	acttogaatt	tctgggcaac	ctgtccggag	atccttatat	tttgac	417

<210> 849  
 <211> 291  
 <212> DNA  
 <213> Pinus radiata

<400> 849

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tcatattctc	tgtagacgag	actcccgttc	gagtgtttta	gaacagggag	acagagttgg	180
gtaaagtgga	tagcaattat	cactatccta	agagccaagc	aatgaaggtc	tattcaagcc	240
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<210> 850  
 <211> 299  
 <212> DNA  
 <213> Pinus radiata

<400> 850

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ctctgctggc	accggtattg	ctttctatat	gtcttcccaa	ggggacgaac	acgatgaatt	180
tgactttgaa	tttttggtta	acatttctgg	acagccatac	actgtgcaga	ccaatgttta	240
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<210> 851  
 <211> 359  
 <212> DNA  
 <213> Pinus radiata

<400> 851

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aatggccaag	tatggcacct	ggttcttgac	cagaactcag	gttctgggtt	caagtccaag	180
tataagtaca	gattcggatg	gttttagcatg	aagctcaagc	tcgtaccggg	agactctgca	240
ggagttgtca	ctgcttatta	tatgtcttct	aacaccgaca	tgaataggga	cgagctggac	300
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<210> 852  
 <211> 347  
 <212> DNA  
 <213> Pinus radiata



&lt;400&gt; 852

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ccctcgggtg gccagttat ccgcattcca	caggctcgaa tagatcttca ttggttggtt	120
gaaagggaac cttatcccca ggttcttgca	gttcttgaa acgcggattg gtaccgagtc	180
cacaaaaaaa ctgcacaaaa aacccatcac	acacatcaat attcatgaac gttccagtgc	240
caaattttga aaagtgtttt tgggttttta	ggattttatt tgtttagaat tttggaagga	300
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&lt;210&gt; 853

&lt;211&gt; 434

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 853

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&lt;210&gt; 854

&lt;211&gt; 274

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 854

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&lt;210&gt; 855

&lt;211&gt; 366

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 855

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&lt;210&gt; 856

&lt;211&gt; 398

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 856

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&lt;210&gt; 857

&lt;211&gt; 183

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 857

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&lt;210&gt; 858

&lt;211&gt; 464

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 858

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&lt;210&gt; 859

&lt;211&gt; 412

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 859

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&lt;210&gt; 860

&lt;211&gt; 376

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 860

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&lt;210&gt; 861

&lt;211&gt; 536

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 861

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&lt;210&gt; 862

&lt;211&gt; 358

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 862

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&lt;210&gt; 863

&lt;211&gt; 322

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 863

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&lt;210&gt; 864

&lt;211&gt; 372

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 864

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<212> DNA  
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<211> 605  
<212> DNA  
<213> Pinus radiata

<400> 868  
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gacct						605

&lt;210&gt; 869

&lt;211&gt; 528

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 869

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&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 870

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&lt;210&gt; 871

&lt;211&gt; 501

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 871

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&lt;210&gt; 872

&lt;211&gt; 540

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 872

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&lt;210&gt; 873

&lt;211&gt; 397

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 873

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&lt;211&gt; 371

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 874

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&lt;211&gt; 355

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 875

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&lt;210&gt; 876

&lt;211&gt; 337

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 876

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&lt;210&gt; 877

&lt;211&gt; 558

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 877

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&lt;210&gt; 878

&lt;211&gt; 400

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 878

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&lt;211&gt; 500

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 879

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&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 880

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&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 881

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&lt;211&gt; 622

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 882

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&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 883

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&lt;212&gt; DNA

&lt;213&gt; Pinus radiata



&lt;400&gt; 884

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&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 885

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&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 886

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&lt;211&gt; 343

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 887

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&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 888

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&lt;211&gt; 543

&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 889

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&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 890

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&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

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&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

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&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

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&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 898

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&lt;212&gt; DNA

&lt;213&gt; Pinus radiata

&lt;400&gt; 900

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ctttgatgca aaactgagga gggaaagtgt agatgtagaa gatctgttca ttgctcatat	180					
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 cgtcctctgt ggggaactgc ccagttgttt atgcatcccc gctacatgca tggagctgct 180  
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## INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

<b>(51) International Patent Classification <sup>7</sup> :</b> C12N 15/53, 15/54, 15/56, 15/60, 15/61, 9/08, 9/10, 9/12, 9/16, 9/18, 9/26, 9/32, 9/38, 9/40, 9/44, 9/88, 9/90, C07K 14/415, A01H 1/00, 5/00	<b>A3</b>	<b>(11) International Publication Number:</b> <b>WO 00/22092</b>  <b>(43) International Publication Date:</b> 20 April 2000 (20.04.00)
<b>(21) International Application Number:</b> PCT/NZ99/00169  <b>(22) International Filing Date:</b> 8 October 1999 (08.10.99)  <b>(30) Priority Data:</b> 09/170,862 13 October 1998 (13.10.98) US 60/148,426 11 August 1999 (11.08.99) US  <b>(71) Applicants (for all designated States except US):</b> GENESIS RE- SEARCH AND DEVELOPMENT CORPORATION LIM- ITED [NZ/NZ]; 1 Fox Street, Parnell, Auckland (NZ). FLETCHER CHALLENGE FORESTS LIMITED [NZ/NZ]; 585 Great South Road, Penrose, Auckland (NZ).  <b>(72) Inventor; and</b> <b>(75) Inventor/Applicant (for US only):</b> BLOKSBERG, Leonard, Nathan [US/NZ]; 5A Korau Road, Greenlane, Auckland (NZ).  <b>(74) Agents:</b> BENNETT, Michael, Roy et al.; West-Walker Ben- nett, Mobil on the Park, 157 Lambton Quay, Wellington (NZ).		<b>(81) Designated States:</b> AE, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CR, CU, CZ, DE, DK, DM, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, US, UZ, VN, YU, ZA, ZW, ARIPO patent (GH, GM, KE, LS, MW, SD, SL, SZ, TZ, UG, ZW), Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European patent (AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE), OAPI patent (BF, BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG).  <b>Published</b> <i>With international search report.</i>  <b>(88) Date of publication of the international search report:</b> 13 July 2000 (13.07.00)
<b>(54) Title:</b> MATERIALS AND METHODS FOR THE MODIFICATION OF PLANT CELL WALL POLYSACCHARIDES		
<b>(57) Abstract</b>  Novel isolated polynucleotides and polypeptides associated with the synthesis of plant cell wall polysaccharides are provided, together with genetic constructs comprising such sequences. Methods for using such constructs for the modulation of polysaccharide content in plants are also disclosed, together with transgenic plants comprising such constructs.		



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EE	Estonia						

# INTERNATIONAL SEARCH REPORT

International application No.

PCT/NZ99/00169

<b>A. CLASSIFICATION OF SUBJECT MATTER</b>		
Int. Cl. <sup>7</sup> : C12N 15/53, 15/54, 15/56, 15/60, 15/61, 9/08, 9/10, 9/12, 9/16, 9/18, 9/26, 9/32, 9/38, 9/40, 9/44, 9/88, 9/90; CO7K 14/415; A01H 1/00, 5/00.		
According to International Patent Classification (IPC) or to both national classification and IPC		
<b>B. FIELDS SEARCHED</b>		
Minimum documentation searched (classification system followed by classification symbols) SEE ELECTRONIC DATABASE BOX BELOW.		
Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched SEE ELECTRONIC DATABASE BOX BELOW.		
Electronic data base consulted during the international search (name of data base and, where practicable, search terms used) EMBL, GENBANK, SWISS PROTEINS, PIR as per sequence id nos specified in inventions 1-5 stated on the extra sheets (1) and (2)..		
<b>C. DOCUMENTS CONSIDERED TO BE RELEVANT</b>		
Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	EP 137280 A (CETUS CORPORATION) 17 April 1985. See the sequence in table 1 on pages 38-44.	1-5, 18 and 19. (Seq id no 125)
X	Sims P F G <i>et al</i> "Differential expression of multiple exocellobiohydrolase I-like genes in the lignin-degrading fungus <i>Phanerochaete chrysosporium</i> " Mol. Microbiol (1994) 12(2) 209-216. See figures 2 and 3 on pages 211 and 213.	1-9, 18 and 19. (seq id nos 129 and 130)
X	Genepept acc. no.. AAB37767 (5 December 1996) See the whole abstract.	1-5, 18 and 19. (seq id nos 38, 40-43, 90, 132-134 and 146.)
<input checked="" type="checkbox"/> Further documents are listed in the continuation of Box C <input checked="" type="checkbox"/> See patent family annex		
* Special categories of cited documents: "A" document defining the general state of the art which is not considered to be of particular relevance "E" earlier application or patent but published on or after the international filing date "L" document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified) "O" document referring to an oral disclosure, use, exhibition or other means "P" document published prior to the international filing date but later than the priority date claimed "T" later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention "X" document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone "Y" document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art "&" document member of the same patent family		
Date of the actual completion of the international search 11 April 2000		Date of mailing of the international search report 14 APR 2000
Name and mailing address of the ISA/AU AUSTRALIAN PATENT OFFICE PO BOX 200, WODEN ACT 2606, AUSTRALIA E-mail address: pct@ipaustalia.gov.au Facsimile No. (02) 6285 3929		Authorized officer J.H. CHAN Telephone No : (02) 6283 2340

# INTERNATIONAL SEARCH REPORT

International application No.

PCT/NZ99/00169

C (Continuation). DOCUMENTS CONSIDERED TO BE RELEVANT		
Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	Genpept acc. No. AAC29067 (7 August 1998) See the whole abstract.	1-5, 18 and 19. (seq id nos 39, 132, 136 and 138)
X	Genpept acc. no. AAC39333 (4 February 1998) See the whole abstract.	1-5, 18 and 19. (seq id nos 39, 41, 93 and 137)
X	Genpept acc. no. AAC39336 (6 February 1998) See whole abstract.	1-5, 18 and 19. (seq id nos 37, 38, 41-43, 93 and 146)
X	EMBL acc. no. Z22528 (8 February 1996) See the whole abstract	1-5, 18 and 19. (Seq id nos 120 and 125)
X	Genbank acc. no. AF27173 (7 February 1998) See whole abstract.	1-5, 18 and 19. (seq id nos 10, 12, 122, 123, 126, 128 and 141)
X	Genbank acc. no. AF27174 (7 February 1998) See whole abstract.	1-5, 18 and 19. (seq id nos 1, 12-14, 70, 122, 124, 128 and 141)
X	Genbank acc. no. AF030052 (5 February 1998) See the whole abstract.	1-5, 18 and 19. (seq id nos 12, 67 and 70)
X	Genbank acc. no. AF062485 (8 August 1998) See the whole abstract.	1-5, 18 and 19. (seq id nos 70, 122, 128 and 144)
X	Genbank acc. no. U58283 (5 December 1996) See the whole abstract.	1-5, 18 and 19. (seq id nos 11, 12, 14, 69, 107, 121, 123, 128 and 141)
X	Genbank acc. no. U58284 (5 December 1996) See the whole abstract.	1-5, 18 and 19. (seq id nos 8, 9, 11, 14, 66, 123, 124 and 128)
X Y	WO 94/28146 A (HOECHST SCHERING AGREEVO GmbH) 8 December 1994. See whole document especially sequence id nos 2 and 4.	1-9, 11-13, and 15-19. (seq id nos 2, 15-17, 19, 59-61, 71-76, 139, and 552-554)

# INTERNATIONAL SEARCH REPORT

International application No.  
PCT/NZ99/00169

C (Continuation) DOCUMENTS CONSIDERED TO BE RELEVANT		
Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	US 5773693 A (Diane G BURGESS and Hugo K DOONER) 30 June 1998. See the whole document, especially sequence id no 5 on columns 41-44.	1-9, 11-13 and 15-19. (seq id no 60)
X	Genpept acc. no. AAB40723 (15 January 1997) See the whole abstract.	1-5, 18 and 19. (seq id nos 32, 87 and 147)
X	Genpept acc. no. AAC49941 (9 March 1998) See the whole abstract.	1-5, 18 and 19. (seq id nos 32 and 38)
X	Genpept acc. no. AAA66057 (10 May 1995) See the whole abstract.	1-5, 18 and 19. (seq id nos 30, 31 and 83-85)
X	Genpept acc. no. AF068260 (11 June 1998) See the whole abstract.	1-5, 18 and 19. (seq id no 65)
X	Swiss-prot acc. no. P55232 (1 October 1996) See the whole abstract.	1-5, 18 and 19. (seq id nos 83, 85 and 87)
X	Swiss-prot acc. no. Q00081 (1 April 1993) See the whole abstract.	1-5, 18 and 19. (seq id nos 32, 88 and 147)
X	Genbank acc. no. U81033 (15 January 1997) See the whole abstract.	1-5, 18 and 19. (seq id nos 3 and 142)
X	Genbank acc. no. U85496 (9 March 1998) See the whole abstract.	1-5, 18 and 19. (seq id no 63)
X	Genbank acc. no. X96765 (13 March 1997) See the whole abstract.	1-5, 18 and 19. (seq id nos 1, 59 and 64)
X Y	WO 96/12814 A (DANISCO A/S) 2 May 1996. See the whole document, especially sequence id no 18	1-9, 11-13 and 15-19. (seq id nos 4-6, 57 and 140)
X Y	US 5316931 A (Jon DONSON <i>et al</i> ) 31 May 1994. See the whole document, especially sequence id no 5 on columns 49-54.	1-9, 11, 12, 18 and 19. (seq id no 58)

# INTERNATIONAL SEARCH REPORT

International Application No.  
PCT/NZ99/00169

C (Continuation) DOCUMENTS CONSIDERED TO BE RELEVANT		
Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X Y	WO 90/12876 A (AKTIESELSKABET DE DANSKE SPRITFABRIKKER) 1 November 1990. See whole document especially figures 2-5.	1-9, 11-13, 15-19. (Seq id nos 34, 35, 81 and 145)
X Y	US 5688684 A (Naoshiro Yoshigi <i>et al</i> ) 18 November 1997. See the whole document, especially sequence id nos 2 and 3 on columns 9-16.	1-9, 11, 12, 18 and 19. 13 and 15-17 (seq id nos 254, 256, and 260-262)
X	Mori H <i>et al</i> "Moklecular cloning of an $\alpha$ - amylase cDNA from germinating cotyledons of kidney bean" J Appl. Glycosience 45(3) 261-267 (1998) See the whole document, especially the amino acid and nucleotide sequences depicted in figure 3 on page 264.	1-5, 18 and 19. (seq id nos 33, 82 and 145)
X	Genpept acc. no. AAA32935 (27 April 1993) See the whole abstract.	1-5, 18 and 19 (seq id no 34)
X	Swiss-Prot acc. no. P17859 (1 August 1990) See the whole abstract.	1-5, 18 and 19. (seq id nos 33 and 82)
X	Embl acc. no. M92090 (27 April 1993) See the whole abstract.	1-5, 18 and 19. (seq id no 253)
X	Embl acc. no. D01022(18 June 1993) See the whole abstract.	1-5, 18 and 19. (seq id nos 252, 254 and 255)
X	Genbank acc. no. Z25871 (6 September 1993) See the whole abstract.	1-5, 18 and 19. (seq id no 261)
X	Genbank acc. no. AJ225087 (25 March 1998) See the whole abstract.	1-5, 18 and 19. (seq id nos 257, 259, 260 and 262)
X	WO 97/32027 A (MAX-PLANCK-GESELLSCHAFT ZUR FORDERUNG DER WISSENSCHAFTEN EV) 4 September 1997. See the whole document, especially sequence id nos 1 and 2 on pages 17-18.	1-10, 11-13 and 15-19. (seq id nos 77 and 78)
X	Hesse H Willmitzer L "Expression analysis of a sucrose synthase gene from sugar beet" Plant Mol. Biol. 30 863-872 (1996) See the whole document, especially the sequences depicted in figure 1 on page 866.	1-10, 11-13 and 15-19. (seq id nos 45, 95-99)

# INTERNATIONAL SEARCH REPORT

International Application No.  
PCT/NZ99/00169

C (Continuation) DOCUMENTS CONSIDERED TO BE RELEVANT		
Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	Godt D E et al "Regulation of sucrose synthase expression on chenopodium rubrum characterization of sugar induced expression in photoautotropho suspension cultures and sink specific expression in plants" J Plant Physiol 146 231-238 (1995) See the whole document, especially the sequence depicted in figure 1 on page 234.	1-9, 11-13, 15-19. (seq id no 102)
X	Genbank acc. no. AF030231 (21 January 1998) See the whole abstract.	1-5, 18 and 19. (seq id nos 15, 16, 18, 105 and 139)
X	PIR acc. no. S22535 (3 May 1994) See the whole abstract.	1-5, 18 and 19. (seq id no 148)
X	Swiss-prot acc. no. Q00917 (1 April 1993) See the whole abstract.	1-5, 18 and 19. (seq id nos 45, 46, 95-97 and 100)
X	Swiss-prot acc. no. Q01390 (1 April 1993) See the whole abstract.	1-5, 18 and 19. (seq id nos 44, 48, 100, 106 and 144)
X	Swiss-prot acc. no. P13708 (1 January 1990) See the whole abstract.	1-5, 18 and 19. (seq id nos 44, 46-48, 98-100, 106 and 144)
X	Swiss-prot acc. no. P31926 (1 July 1993) See the whole abstract.	1-5, 18 and 19. (seq id nos 44, 47, 100, 101, 106 and 144)
P, X	WO 98/53085 A (ZENECA LIMITED) 26 November 1998 (Priority date 20 May 1997) See the whole document, especially sequence id no 42 on page 42-43.	1-11, 18 and 19. (seq id nos 20, 21, 79 and 80)
X	Swiss-prot acc. no. P19595 (1 February 1991) See the whole abstract.	1-5, 18 and 19. (seq id nos 49, 50, 51, 103 and 104)
X	Genbank acc. no. Z18924 (4 December 1992) See the whole abstract.; seq id nos 20, 21, 22, 79 and 80.	1-5, 18 and 19. (seq id nos 20-22, 79 and 80)

# INTERNATIONAL SEARCH REPORT

International application No.

PCT/NZ99/00169

## Box I Observations where certain claims were found unsearchable (Continuation of item 2 of first sheet)

This international search report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:

1. ☐ Claims Nos :  
because they relate to subject matter not required to be searched by this Authority, namely:
  
2. ☐ Claims Nos :  
because they relate to parts of the international application that do not comply with the prescribed requirements to such an extent that no meaningful international search can be carried out, specifically:
  
3. ☐ Claims Nos :  
because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a)

## Box II Observations where unity of invention is lacking (Continuation of item 3 of first sheet)

This International Searching Authority found multiple inventions in this international application, as follows:

The International Searching Authority has found that there are 34 separate inventions, wherein a single enzyme type or protein provides the special technical feature. This is based on the following reasons:

- 1) The international application has claimed nucleic acid sequences encoding 33 different enzymes and 1 protein (annexin), the fragments of these genes, their use in transforming plants to modulate the polysaccharides content of ( to be continued on the extra sheets (1)-(5))

1. ☐ As all required additional search fees were timely paid by the applicant, this international search report covers all searchable claims
2. ☐ As all searchable claims could be searched without effort justifying an additional fee, this Authority did not invite payment of any additional fee.
3. ☒ As only some of the required additional search fees were timely paid by the applicant, this international search report covers only those claims for which fees were paid, specifically claims Nos. 1-19 in the inventions as defined in the following::  
Inventions 1, 2, 3, 4 and 5 as stated on the extra sheets (1) and (2).
4. ☐ No required additional search fees were timely paid by the applicant. Consequently, this international search report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:

### Remark on Protest

- ☐ The additional search fees were accompanied by the applicant's protest.
- ☒ No protest accompanied the payment of additional search fees.

## INTERNATIONAL SEARCH REPORT

International application No.

PCT/NZ99/00169

### Supplemental Box

(To be used when the space in any of Boxes I to VIII is not sufficient)

#### Continuation of Box No II:

the plant, the probes or primers based on these genes, and polypeptides coded by these genes and their variants with at least 50% homology.

2) Whilst these 908 sequences are from either *Pinus radiata* or *Eucalyptus grandis*, the invention is to these sequences *per se* and their variant which have 50% homology to the former. The invention as described is also to the use of oligonucleotide probes or primers based on these sequences. It is clear that all these nucleic acids sequences are not limited to the source from which they are isolated, as such the source from these plants cannot be the special technical feature under Rule 13.2 of the PCT. (3)

3) The nucleic acid sequences and their putative amino acid sequences have been shown to have similarity to protein or enzymes which are known to be involved in the synthesis of polysaccharides in the cell walls of plants (p 9 lines 20 to page 11 lines 2). Based on this methodology, the 909 sequences in the quoted passage have been assigned with 33 different enzymic activities and the biological activity of the annexin (a non-enzymic protein). However, these enzymes and protein are not unified by sequence homology, by a common substrate or their mode of action (eg as a non-enzyme, a hydrolase or an oxido-reductase etc.). In addition, many of these enzymes are known to have activity inside the cell and not associated directly with the syntheses of cell wall polysaccharides. Furthermore, the use of the sense or antisense constructs coding the enzymes as transgenes in plants are known. Several transgenes (eg based on branching enzyme, sucrose synthase, ADP-glucose pyrophosphorylase, UDP-glucose pyrophosphorylase and 1,3-  $\beta$ -glucanase) have been used to alter the polysaccharide contents in the plant or plant cells. Therefore, the use of the nucleotide sequences encoding these proteins as transgenes, either in the sense or antisense direction, to affect polysaccharide content or composition in plant is not a special technical feature under Rule 13.2 of the PCT.

For these reasons the international searching authority has identified 34 inventions; they are as listed below:

1. Nucleic and amino acid sequences SEQ ID NOS 7-14, 36-43, 66-70, 90-94, 107, 108, 119-138, 141, 146 and their at least 50% homologues coding cellulose synthase, DNA probes or primers therefrom, composition containing these nucleic acids, their sense or antisense recombinant constructs, transgenic plant/plant cell based on these constructs and a method of modulating the polysaccharide content of plant or the activities of these polypeptides in the biosynthetic pathway using these constructs.
2. Nucleic and amino acid sequences SEQ ID NOS 1-3, 30-32, 59-65, 83-89, 142, 147 and their at least 50% homologues coding ADP-glucose pyrophosphorylase, DNA probes or primers therefrom, composition containing these nucleic acids, their sense or antisense recombinant constructs, transgenic plant/plant cell based on these constructs and a method of modulating the polysaccharide content of plant or the activities of these polypeptides in the biosynthetic pathway using these constructs.
3. Nucleic and amino acid sequences SEQ ID NOS 4-6, 33-35, 57, 58, 81, 82, 140, 145 and 252-262 and their at least 50% homologues coding amylase,  $\alpha$ - or  $\beta$ -amylase, DNA probes or primers therefrom, composition containing these nucleic acids, their sense or antisense recombinant constructs, transgenic plant/plant cell based on these constructs and a method of modulating the polysaccharide content of plant or the activities of these polypeptides in the biosynthetic pathway using these constructs.
4. Nucleic and amino acid sequences SEQ ID NOS 15-19, 44-48, 71-78, 95-102, 105, 106, 139, 143, 144, 148 and 552-555 and their at least 50% homologues coding sucrose synthase, DNA probes or primers therefrom, composition containing these nucleic acids, their sense or antisense recombinant constructs, transgenic plant/plant cell based on

- to be continued on the extra sheet (2) -



## INTERNATIONAL SEARCH REPORT

International application No.

PCT/NZ99/00169

### Supplemental Box

(To be used when the space in any of Boxes I to VIII is not sufficient)

#### Continuation of Box No II (continued from extra sheet (1))

these constructs and a method of modulating the polysaccharide content of plant or the activities of these polypeptides in the biosynthetic pathway using these constructs.

5. Nucleic and amino acid sequences SEQ ID NOS 20-23, 49-51, 79, 80, 103, 104 and their at least 50% homologues coding UDP-glucose pyrophosphorylase, DNA probes or primers therefrom, composition containing these nucleic acids, their sense or antisense recombinant constructs, transgenic plant/plant cell based on these constructs and a method of modulating the polysaccharide content of plant or the activities of these polypeptides in the biosynthetic pathway using these constructs.
6. Nucleic and amino acid sequences SEQ ID NOS 24-29, 52-56, 109-118 and their at least 50% homologues coding annexin, DNA probes or primers therefrom, composition containing these nucleic acids, their sense or antisense recombinant constructs, transgenic plant/plant cell based on these constructs and a method of modulating the polysaccharide content of plant or the activities of these polypeptides in the biosynthetic pathway using these constructs.
7. Nucleic and amino acid sequences SEQ ID NOS 149-185 and their at least 50% homologues coding 1, 3- $\beta$ -D-glucanase, DNA probes or primers therefrom, composition containing these nucleic acids, their sense or antisense recombinant constructs, transgenic plant/plant cell based on these constructs and a method of modulating the polysaccharide content of plant or the polypeptides in the activities of these biosynthetic pathway using these constructs.
8. Nucleic and amino acid sequences SEQ ID NOS 186 and their at least 50% homologues coding 1,4- $\beta$ -cellobiohydrolase, DNA probes or primers therefrom, composition containing these nucleic acids, their sense or antisense recombinant constructs, transgenic plant/plant cell based on these constructs and a method of modulating the polysaccharide content of plant or the activities of these polypeptides in the biosynthetic pathway using these constructs.
9. Nucleic and amino acid sequences SEQ ID NOS 187-196 and their at least 50% homologues coding  $\alpha$ , $\alpha$ -trehalose phosphate synthase, DNA probes or primers therefrom, composition containing these nucleic acids, their sense or antisense recombinant constructs, transgenic plant/plant cell based on these constructs and a method of modulating the polysaccharide content of plant or the activities of these polypeptides in the biosynthetic pathway using these constructs.
10. Nucleic and amino acid sequences SEQ ID NOS 197-204 and their at least 50% homologues coding  $\alpha$ -glucosidase, DNA probes or primers therefrom, composition containing these nucleic acids, their sense or antisense recombinant constructs, transgenic plant/plant cell based on these constructs and a method of modulating the polysaccharide content of plant or the activities of these polypeptides in the biosynthetic pathway using these constructs.
11. Nucleic and amino acid sequences SEQ ID NOS 205-250 and their at least 50% homologues coding aldolase, DNA probes or primers therefrom, composition containing these nucleic acids, their sense or antisense recombinant constructs, transgenic plant/plant cell based on these constructs and a method of modulating the polysaccharide content of plant or the activities of these polypeptides in the biosynthetic pathway using these constructs.

- to be continued on the extra sheet (3) -

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### Supplemental Box

(To be used when the space in any of Boxes I to VIII is not sufficient)

Continuation of Box No II (continued from extra sheet (2)):

12. Nucleic and amino acid sequences SEQ ID NOS 251 and their at least 50% homologues coding amylopectin 6-glucanohydrolase, DNA probes or primers therefrom, composition containing these nucleic acids, their sense or antisense recombinant constructs, transgenic plant/plant cell based on these constructs and a method of modulating the polysaccharide content of plant or the activities of these polypeptides in the biosynthetic pathway using these constructs.
13. Nucleic and amino acid sequences SEQ ID NOS 263 and their at least 50% homologues coding P-glucosidase, DNA probes or primers therefrom, composition containing these nucleic acids, their sense or antisense recombinant constructs, transgenic plant/plant cell based on these constructs and a method of modulating the polysaccharide content of plant or the activities of these polypeptides in the biosynthetic pathway using these constructs.
14. Nucleic and amino acid sequences SEQ ID NOS 264-272 and their at least 50% homologues coding branching enzyme, DNA probes or primers therefrom, composition containing these nucleic acids, their sense or antisense recombinant constructs, transgenic plant/plant cell based on these constructs and a method of modulating the polysaccharide content of plant or the activities of these polypeptides in the biosynthetic pathway using these constructs.
15. Nucleic and amino acid sequences SEQ ID NOS 273-318 and their at least 50% homologues coding D-fructokinase, DNA probes or primers therefrom, composition containing these nucleic acids, their sense or antisense recombinant constructs, transgenic plant/plant cell based on these constructs and a method of modulating the polysaccharide content of plant or the activities of these polypeptides in the biosynthetic pathway using these constructs.
16. Nucleic and amino acid sequences SEQ ID NOS 319-354 and their at least 50% homologues coding D-xylulose reductase, DNA probes or primers therefrom, composition containing these nucleic acids, their sense or antisense recombinant constructs, transgenic plant/plant cell based on these constructs and a method of modulating the polysaccharide content of plant or the activities of these polypeptides in the biosynthetic pathway using these constructs.
17. Nucleic and amino acid sequences SEQ ID NOS 355-365 and their at least 50% homologues coding endo-1, 3-1,4-p-glucanase, DNA probes or primers therefrom, composition containing these nucleic acids, their sense or antisense recombinant constructs, transgenic plant/plant cell based on these constructs and a method of modulating the polysaccharide content of plant or the activities of these polypeptides in the biosynthetic pathway using these constructs.
18. Nucleic and amino acid sequences SEQ ID NOS 366-371 and their at least 50% homologues coding glucan exo-1, 3-P-glucosidase, DNA probes or primers therefrom, composition containing these nucleic acids, their sense or antisense recombinant constructs, transgenic plant/plant cell based on these constructs and a method of modulating the polysaccharide content of plant or the activities of these polypeptides in the biosynthetic pathway using these constructs.
19. Nucleic and amino acid sequences SEQ ID NOS 372-377 and their at least 50% homologues coding glucose 6-phosphate dehydrogenase, DNA probes or primers therefrom, composition containing these nucleic acids, their sense or antisense recombinant constructs, transgenic plant/plant cell based on these constructs and a method of modulating the polysaccharide content of plant or the activities of these polypeptides in the biosynthetic pathway using these constructs

-to be continued on extra sheet (4)-

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### Supplemental Box

(To be used when the space in any of Boxes I to VIII is not sufficient)

Continuation of Box No II (continued from extra sheet (3)):

20. Nucleic and amino acid sequences SEQ ID NOS 378-38 land their at least 50% homologues coding glucose phosphate isomerase, DNA probes or primers therefrom, composition containing these nucleic acids, their sense or antisense recombinant constructs, transgenic plant/plant cell based on these constructs and a method of modulating the polysaccharide content of plant or the activities of these polypeptides in the biosynthetic pathway using these constructs.
21. Nucleic and amino acid sequences SEQ ID NOS 382-389 and their at least 50% homologues coding isoamylase, DNA probes or primers therefrom, composition containing these nucleic acids, their sense or antisense recombinant constructs, transgenic plant/plant cell based on these constructs and a method of modulating the polysaccharide content of plant or the activities of these polypeptides in the biosynthetic pathway using these constructs.
22. Nucleic and amino acid sequences SEQ ID NOS 390-393 and their at least 50% homologues coding L-ribulokinase, DNA probes or primers therefrom, composition containing these nucleic acids, their sense or antisense recombinant constructs, transgenic plant/plant cell based on these constructs and a method of modulating the polysaccharide content of plant or the activities of these polypeptides in the biosynthetic pathway using these constructs.
23. NucleicandaminoacidsequencesSEQIDNOS394-398andtheiratleast50%homologuescoding mannitol-1-phosphate 5-dehydrogenase, DNA probes or primers therefrom, composition containing these nucleic acids, their sense or antisense recombinant constructs, transgenic plant/plant cell based on these constructs and a method of modulating the polysaccharide content of plant or the activities of these polypeptides 'in the biosynthetic pathway using these constructs.
24. Nucleic and amino acid sequences SEQ ID NOS 399-478 and their at least 50% homologues coding pectin methyl-esterase, DNA probes or primers therefrom, composition containing these nucleic acids, their sense or antisense recombinant constructs, transgenic plant/plant cell based on these constructs and a method of modulating the polysaccharide content of plant or the activities of these polypeptides in the biosynthetic pathway using these constructs.
25. Nucleic and amino acid sequences SEQ ID NOS 479-506 and their at least 50% homologues coding phosphoglucomutase, DNA probes or primers therefrom, composition containing these nucleic acids, their sense or antisense recombinant constructs, transgenic plant/plant cell based on these constructs and a method of modulating the polysaccharide content of plant or the activities of these polypeptides 'in the biosynthetic pathway using these constructs.
26. Nucleic and amino acid sequences SEQ ID NOS 507-508 and their at least 50% homologues coding phospho-ribulokinase, DNA probes or primers therefrom, composition containing these nucleic acids, their sense or antisense recombinant constructs, transgenic plant/plant cell based on these constructs and a method of modulating the polysaccharide content of plant or the polypeptides in the biosynthetic pathway using these constructs.
27. Nucleic and amino acid sequences SEQ ID NOS 509-521 and their at least 50% homologues coding ribulose-phosphate-3-epimerase, DNA probes or primers therefrom, composition containing these nucleic acids, their sense or antisense recombinant constructs, transgenic plant/plant cell based on these constructs and a method of modulating the polysaccharide content of plant or the polypeptides in the biosynthetic pathway using these constructs.

- to be continued on the extra sheet (5) -

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### Supplemental Box

(To be used when the space in any of Boxes I to VIII is not sufficient)

Continuation of Box No. II (continues from extra sheet (4)):

28. Nucleic and amino acid sequences SEQ ID NOS 522-530 and their at least 50% homologues coding starch phosphorylase, DNA probes or primers therefrom, composition containing these nucleic acids, their sense or antisense recombinant constructs, transgenic plant/plant cell based on these constructs and a method of modulating the polysaccharide content of plant or the activities of these polypeptides in the biosynthetic pathway using these constructs.
29. Nucleic and amino acid sequences SEQ ID NOS 531-551 and their at least 50% homologues coding sucrose phosphate synthase, DNA probes or primers therefrom, composition containing these nucleic acids, their sense or antisense recombinant constructs, transgenic plant/plant cell based on these constructs and a method of modulating the polysaccharide content of plant or the activities of these polypeptides in the biosynthetic pathway using these constructs 30. Nucleic and amino acid sequences SEQ ID NOS 556-586 and their at least 50% homologues coding transketolase, DNA probes or primers therefrom, composition containing these nucleic acids, their sense or antisense recombinant constructs, transgenic plant/plant cell based on these constructs and a method of modulating the polysaccharide content of plant or the activities of these polypeptides in the biosynthetic pathway using these constructs.
30. Nucleic and amino acid sequences SEQ ID NOS 556-586 and their at least 50% homologues coding transketolase, DNA probes or primers therefrom, composition containing these nucleic acids, their sense or antisense recombinant constructs, transgenic plant/plant cell based on these constructs and a method of modulating the polysaccharide content of plant or the activities of these polypeptides in the biosynthetic pathway using these constructs.
31. Nucleic and amino acid sequences SEQ ID NOS 587-591 and their at least 50% homologues coding trehalase, DNA probes or primers therefrom, composition containing these nucleic acids, their sense or antisense recombinant constructs, transgenic plant/plant cell based on these constructs and a method of modulating the polysaccharide content of plant or the activities of these polypeptides in the biosynthetic pathway using these constructs.
32. Nucleic and amino acid sequences SEQ ID NOS 592-620 and their at least 50% homologues coding UDP-glucose 4-epimerase, DNA probes or primers therefrom, composition containing these nucleic acids, their sense or antisense recombinant constructs, transgenic plant/plant cell based on these constructs and a method of modulating the polysaccharide content of plant or the activities of these polypeptides in the biosynthetic pathway using these constructs.
33. Nucleic and amino acid sequences SEQ ID NOS 621-902 and their at least 50% homologues coding xyloglucan endo transglycosylase, DNA probes or primers therefrom, composition containing these nucleic acids, their sense or antisense recombinant constructs, transgenic plant/plant cell based on these constructs and a method of modulating the polysaccharide content of plant or the activities of these polypeptides in the biosynthetic pathway using these constructs.
34. Nucleic and amino acid sequences SEQ ID NOS 903-908 and their at least 50% homologues coding xylose isomerase, DNA probes or primers therefrom, composition containing these nucleic acids, their sense or antisense recombinant constructs, transgenic plant/plant cell based on these constructs and a method of modulating the polysaccharide content of plant or the activities of these polypeptides in the biosynthetic pathway using these constructs

# INTERNATIONAL SEARCH REPORT

Information on patent family members

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This Annex lists the known "A" publication level patent family members relating to the patent documents cited in the above-mentioned international search report. The Australian Patent Office is in no way liable for these particulars which are merely given for the purpose of information.

Patent Document Cited in Search Report				Patent Family Member			
EP	137280	AU	32530/84	BR	8404346	CA	1338400
		ES	535511	ES	543766	ES	543767
		ES	8602139	ES	8604304	ES	860305
		JP	60149387	JP	7051071		
WO	9428146	DE	4317596	EP	701617	HU	74394
		US	5866790				
US	5773693	US	5498831				
WO	9612814	AU	27881/95	CA	2202896	EP	787194
US	5316931	AU	40725/89	EP	406267	US	5589367
		US	5866785	US	5889190	US	5922602
		US	5529909				
WO	9012876	AU	55318/90	CA	2053230	EP	470145
		US	5498832	US	5789657		
WO	9732027	DE	19607697	EP	883689		
WO	9853085	AU	72257/98				
US	5688684	EP	713916	US	5863784	JP	8089245
END OF ANNEX							